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OM protein - protein search, using sw model
                                                                                                                                                                                                 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 50 summaries
                                                                                                                                                                                                                                                                             Minimum DB seq length: 0
Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Perfect score:
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                                                                                                                                                             Database
                                                                                                                                                                                                                                                                                                                                          Total number of hits satisfying chosen parameters:
                                                                                                  A_Geneseq_16Dec04:*
1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
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32
1 IGDIGD 6
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(without alignments)
14.150 Million cell updates/sec
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geneseqp2001s:*
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geneseqp2003bs:*
geneseqp2004s:*
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                                                                                                                                                                                                                                                                                                                                                                                  386760381 residues
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	80	7	σ	υ	4.	ω	2	سر	Result No.
31	31	31	31	31	31	31	31	31	31	31	31	31	31	31	31	31	31	31	31	31	32	32	32	32	Score
96.9	96.9	9	٠	96.9	٥,	<u>,</u>	96.9	96.9	96.9	96.9	٥.	96.9	٥.	9	٥,	٥.	96.9	96.9	٥,	96.9	100.0	100.0	100.0	100.0	Query Match
364	342	220	207	205	201	173	173	166	166	166	166	166	156	149	149	129	86	81	69	59	515	443	6	6	Length
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ADS45014	ADJ48363	ADQ10053	AAM79529	AAU18546	ADQ10039	ABM48831	AAU52312	ADI24536	ADJ70104	ADB75248	ABR41956	AAM78545	AAU18521	ABM47745	AAU51226	AAU18547	AAG01408	AB055461	AAG01407	AAW22623	ADS29151	ABP65456	ADG43178	3	ID
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Ads45014	Adj48	Adq10053	Aam79529	Aau18546	Adq10039	Abm48831	Aau52312	Ad124536	Adj 70104	Adb75248	Abr41956	Aam78545	Aau18521	Abm47745	Aau51226	Aau18547	Aag01408	Abo55461	Aag01407	Aaw22623	Ads29151	Abp65456	Adg43178	Adg43179	Descr
5014 Bacterial		B_afze		×					Human	Prost	1956 Mouse cof	Human		-			Humar	3461 Human gen	Humar	2623 Type II t	151 Bacterial	456 Bifidobac		IGD-moti	Description

ALIGNMENTS

RESULT 1

ADG4	
Χij	ADG43179 standard; peptide; 6 AA.
AC	ADG43179;
S E	26-FEB-2004 (first entry)
BX	IGD-motif peptide #SEQ ID 4.
žX	Andiogenic composition: isoleucine-glycine-aspartic acid; IGD;
Z Z	migration; angiogenesis; is
Ş	polyvinyl pyrrolidone; cyclic.
2 \$	Thidestified
¥ 8	VIIAUGIICAAAAA
Ng Y	WO2003059436-A2.
PD	24-JUL-2003.
무소	23-DEC-2002; 2002WO-US041484.
PR X	21-DEC-2001; 2001US-00027015.
PX	(CENT-) CENTERPULSE BIOLOGICS INC.
2 2	Akella R, Ranieri J;
g X	WPT: 2003-697294/66.
i X	The important composition comprising pentides containing isolenging of
r d	aspartic acid, that stimulate cell migration, and angiogenic growth
F X	factor useful for promoting myocardial or peripheral anglogenesis.
Sd.	Claim 1; SEQ ID NO 4; 22pp; English.
28	The invention relates to an angiogenic composition (I) comprising
ຄຸຄ	
ဂ္ဂ	factor other than the at least one po
გგ	peptide stimulates cell migration. Also disclosed is a composition that is active for promoting cell migration and/or angiogenesis un
8	growth promoting conditions comprising an IGD

heart;

The invention relates to an angiogenic composition (I) comprising at least one peptide chosen from a group of peptides containing isoleucine-glycine-aspartic acid (IGD-containing peptide), and at least one angiogenic growth factor other than the at least one peptide, where the peptide stimulates cell migration. Also disclosed is a composition (III) that is active for promoting cell migration and/or angiogenesis under cell growth promoting conditions comprising an IGD-containing peptide,

Angiogenic composition comprising peptides containing isoleucine-glycine-aspartic acid, that stimulate cell migration, and angiogenic growth factor useful for promoting myocardial or peripheral angiogenesis.

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RESULT 2
ADG43178
ID ADG4
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The invention relates to an angiogenic composition (I) comprising at least one peptide chosen from a group of peptides containing isoleucine-glycine-aspartic acid (IGD-containing peptide), and at least one angiogenic growth factor other than the at least one peptide, where the peptide stimulates cell migration. Also disclosed is a composition (III) that is active for promoting cell migration and/or angiogenesis under cell growth promoting conditions comprising an IGD-containing peptide, and a matrix material. Compositions of the invention are useful for promoting myocardial angiogenesis which involves administering (I) or (III) intramyocardially to an isschaemic area of the heart of an individual, to enhance vascular endothelial cell migration and/or proliferation. They are also useful for promoting peripheral angiogenesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    migration and/or proliferation surficient to receptable carrier in flow to the ischaemic tissue. The physiologically acceptable carrier in the composition is polyvinyl pyrrolidone. The current sequence represents an IGD-motif peptide that is specifically claimed in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     proliferation. They are also useful for promoting peripheral angiogenesis in an ischaemic region of an organ or tissue fed by a peripheral vessel, by enhancing vascular endothelial cell misration and/or proliferation at the ischaemic region, or for enhancing blood flow to an ischaemic tissue of the body, where the composition stimulates vascular endothelial cell migration and/or proliferation sufficient to restore or increase blood migration and/or proliferation sufficient to restore or increase blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             promoting myocardial angiogenesis which involves administering (I) (III) intramyocardially to an ischaemic area of the heart of an individual, to enhance vascular endothelial cell migration and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Angiogenic composition comprising peptides containing isoleucine-glycine-
aspartic acid, that stimulate cell migration, and angiogenic growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Angiogenic composition; isoleucine-glycine-aspartic acid; IGD; angiogenic growth factor; cell migration; angiogenesis; ischae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG43178 standard; peptide; 6 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 2; SEQ ID NO 3; 22pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      factor useful for promoting myocardial or peripheral angiogenesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-697294/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-DEC-2001; 2001US-00027015
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IGD-motif peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CENT-) CENTERPULSE BIOLOGICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     matrix material. Compositions of the invention are useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 IGDIGD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INC
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Pred. No. 1.8e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 migration; angiogenesis; ischaemia; heart;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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       angiogenesis
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                         sequence of a Bifidobacterium genome selected from the nucleotide (c) sequences given in ABQ81842 and ABQ81843, or a sequence exhibiting at (c) least 90% identity or which hybridises with the sequence exhibiting at (c) least 90% identity or which hybridises with the sequence siven in CC ABQ81843. Also described is a polynucleotide (II) encoding a CC fusion protein, comprising a sequence selected from 1097 sequences given in ABP65258 to ABP66354 ligated in frame to a polynucleotide encoding a cc in the cologous polypeptide. (I) has antidiarrheic and antibacterial (C) activities, and can be used as an inhibitor of Salmonella. (I) (which is a probe) is useful for the detection and/or identification of Bifidobacterium longum in a biological sample. A carrier containing the lactic acid bacterium Bifidobacterium longum NCC2705 (CNCM I-2618) can be used for preventing and/or treating diarrhoea brought about by pathogenic bacteria and/or rotarirus. The carrier is a food composition selected from milk, yogurt, curd, cheese, fermented milks, milk based fermented convolutes, ife-creams, fermented cereal based products, ice-creams, fermented cereal based products.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Jacter yogur. from milk, yogur. re-
rroducts, ice-creams, re-
rroducts 'nfant formula,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel polynucleotide comprising Bifidobacterium genome sequence useful a probe or primer for detecting and/or identifying Bifidobacterium long
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bifidobacterium longum NCC2705; Bifidobacterium; bacterial; antidiarrheic; antibacterial; inhibitor of Salmonella; dete
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bifidobacterium
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                                                                                                                                                                                                                                                                                                                                                                                             Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                in a biological sample.
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                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 200; 80pp; English.
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                                                                                                                                                                                                                                                                                                                           invention describes a polynucleotide (I) comprising a Bifidobacterium genome selected from the nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ilarity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lactic acid bacterium; diarrhoea; pathogenic bacteria; composition; pharmaceutical composition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    longum NCC2705 ORF amino acid sequence
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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Matches
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Best Local
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The invention relates to a recombinant DNA construct comprising a promoter functional in a plant cell, where the promoter is positioned to provide for expression of a polynucleotide encoding a polypeptide from a microbial source. The invention also relates to a transformed plant comprising the recombinant DNA construct and a method of producing a transformed plant having an improved property. The plant is a crop plant such as maize or soybean. The method of producing a transformed plant having an improved property at ransformed plant with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cold tolerance; heat tolerance; drought tolerance; herbicide; osmosis; pathogen tolerance; pest tolerance; plant disease resistance; cell cycle pathway modification; plant growth regulator; homologous recombination; seed oil yield; protein yield; carbohydrate; nitrogen; phosphorus; photosynthesis; lignin; galactomannan; bacterial polypeptide.
                                                                                                                                                                                                                                                                            New recombinant DNA construct comprising a promoter positioned to provide for expression of a polynucleotide encoding a polypeptide from a microbial source, useful for producing plants with improved properties.
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(HINK/)
(SLAT/)
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                                                                                                                                                                                                                                 Claim 1;
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) HINKLE G J.
) SLATER S C.
) CHEN X.
) GOLDMAN B S.
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Pred. No.
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Claim

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RESULT 5
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25-MAR-2003
12-NOV-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Type II topoisomerase database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW22623;
                                                                                                                                                                                  05-JUL-1990;
13-AUG-1993;
                                                                                                                                                                                                                                 06-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ignigh 428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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93US-00106482.
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                                                                                                                                                                                                                                                                                                                                                                                         /note= "Variable segment 2 of type II topoisomerase"
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                                                                                                                                                                                                                                                                                                                                                       note= "Variable segment
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Pred. No.
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                                                                                                                                                                                                                                                                                                                               II topoisomerase*
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CC (Updated on 25-MAR-2003 to correct PF field.) (Updated on 27-AUG-2003 to
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Best Local S
Matches 5
The present sequence is a polypeptide encoded by one of a large number of STSTs derived from mRNAs encoding secreted proteins. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT
                                                                                                                                                                                                                                             New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted protein, SEQ ID NO: 5488.
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                                                                                                                                                                    Claim 13; SEQ ID NO 5488;
                                                                                                                                                                                                                                                                                                                                  N-PSDB;
                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-500381/45.
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                                                                                                                                                                                                                 forensic, gene therapy and chromosome mapping procedures
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Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                  Giordano J;
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                                                                                                                                                                 English.
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RESULT 7
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Best Local
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                                                                                                                                                                                                                           New human genome-derived single exon nucleic acid probes useful for human gene expression analysis, for identifying or characterizing alternative splicing events, for assessing genomic alterations or as tools for
                                                                                                                                                                                                                                                                                                                                                                         03-APR-2002; 2002US-00029386
                                                                                                                                                                                                                                                                                                                                                                                                                        16-OCT-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; gene expression; single exon probe; microarray; alternative splicing event; genomic alteration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human genome derived single exon protein #1695
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABO55461;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 69 AA;
                                                                                                                                                                                                                                                                            WPI; 2004-119264/12.
                                                                                                                                                                                                                                                                                                                                                                                                03-APR-2002; 2002US-00029386
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                         (PENN/) PENN S G.
(RANK/) RANK D R.
(HANZ/) HANZEL D K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  40 VGDIGD 45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 5; Conserv
                                                                                                                                                                                                                                                                                                   Rank DR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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The invention relates to a nucleic acid probe for measuring human gene expression, comprising any of the 27,400 fully defined nucleotide sequences in the specification, or their complements or fragments, and encoding at least 8 amino acids of any of the 6888 amino acid sequences fully defined in the specification. The probe is a single exon probe that hybridises under high stringency conditions to a nucleic acid molecule expressed in human cells or tissues. Also included are a spatially-addressable set of single exon nucleic acid grobes for measuring human gene expression (comprising a plurality of single exon nucleic acid probes is separately and addressably isolatable or amplifiable from the plurality), a single exon microarray for measuring human gene expression, a vector comprising at least 8 more probe cited above, an ORF-encoded peptide comprising at least 8

acids

of.

Claim 45; SEQ ID NO 29095; 80pp; English.

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AAGO1408
ID AAGC
XX AAGC
XX O6-C
XX Huma
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequences (optionally with conservative amino acid substitutions), an isolated antibody that binds specifically to a peptide cited above, methods of selling and/or licensing single exon probes or microarrays to a customer desiring to measure gene expression, a method of providing human gene expression data by subscription, and a computer-readable storage medium which contains a database having a plurality of records (each record including data on the expression of a single exon probe cited above. The probe, methods and apparatus are useful in gene expression analysis. The probes may be used as tools for surveying tissues to detect the presence of expressed messages that contain their confidence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            smaller genomic alterations, in priming the synthesis of nucleic acids, or in expressing the ORF-encoded peptide. The present sequence is a human single exon probe protein of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from USPTO at segdata.uspto.gov/sequence.html?DocID=20030194704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     specific exon, or in constructing genome-derived single exon microarrays. In addition, the probes are used in identifying and characterising alternative splicing events, in detecting and characterising gross alternations in the genomic locus that includes their exon, in assessing alternations in the genomic locus that includes their exon, in assessing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene
                                                                                                                                                                                                                                                                                                              Dumas Milne Edwards
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; 5' EST;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-OCT-2000
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                                                                                                                                                                                       New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo
                                                                                                                                                                                                                                                   N-PSDB; AAC01414
                                                                                                                                                                                                                                                                                                                                                                                            26-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                    21-FEB-2000; 2000EP-00200610
                                                                                                                                                                                                                                                                                                                                                    (GEST ) GENSET
                                                                                                                                                                  diagnostic,
                                                                                                                                                                                     obtaining cDNAs and genomic
                                                                                                                                                                                                                                                                   2000-500381/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   therapy;
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                                                                                                                                                                  forensic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; expressed chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein,
                                                                                                                                                                                                                                                                                                                                                                                              99US-0122487P
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83.3%;
                                                                                                                                                                gene therapy and chromosome
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                                                                                                                                                                                                                                                                                                              Duclert A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence tag; secreted protein; cDNA isolation; mapping.
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Pred. No. 1.6e+02;
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                                                                                                                                                                                                                                                                                                              Giordano
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5' ESTs derived from mRNAs encoding secreted proteins. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated

polypeptide encoded by one of a large number of

region

of the mRNA because

Claim 13;

SEQ ID NO 5489; 71pp + Sequence Listing; English.

The present sequence is a psychology of the present sequence is a psychology of the present and present the presen

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RESULT 9
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Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cytoskeletal element-related protein; human; mouse; rabbit; goat; horse; cat; dog; chicken; sheep; immunosuppressive; antiarthritic; vasotropic; antirheumatic; antiproliferative; cytostatic; cardiant; neuroprotective; cerebroprotective; nootropic; antibacterial; virucide; fungicide; cancer; ophthalmological; vulnerary; gene therapy; autoimmune disease; neoplasm; hyperproliferative disorder; breast; liver; cardiovascular disorder; hyperproliferative disorder; breast; liver; cardiovascular disorder; fungal infection; viral infection; ocular disorder; endocrine disorder; gastrointestinal disorder; renal disorder; respiratory disorder; wound healing; skin aging; organ transplantation; food preservative; tissue regeneration; anti-infertility; food additive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion vectors
                                                                                                                                                                                                                               07-JUL-2000;
11-JUL-2000;
11-JUL-2000;
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02-MAR-2000;
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07-JUL-2000;
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19-MAY-2000;
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17-MAR-2000;
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2000US-0205515P
2000US-0209467P
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2000US-0217496P
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2000US-0224519P
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2000US-0225214P
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83.3%;
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Pred. No. 1.7e+02;
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Sequences AAU18508-AAU18551 represent the cytoskeletal element-related CC polypeptides of the invention. Cytoskeletal polypeptides and their CC associated polynucleotides are useful in the diagnosis, treatment and CC prevention of various types of disorders in e.g. humans, mice, rabbits, CC goats, horses, cats, dogs, chickens or sheep. A pathological condition CC can be determined by determining the presence or absence of a mutation in a cytoskeletal polynucleotide. The treatable disorders include autoimmune CC diseases such as rheumatoid arthritis, hyperproliferative disorders such as cardiaca arrest, cerebrovascular disorders such as crebral ischaemia, CC as neoplasms of the breast or liver, cardiovascular disorders such as caused by bacteria, viruses and fungi, ocular disorders such as corneal CC infection, endocrine disorders such as premature labour and infertility, gastrointestinal disorders such as gromature labour and infertility, as glomerulonephritis and respiratory disorders such as asthma. The CC polypeptides can also be used to aid wound healing, to prevent skin aging due to sunburn, to maintain organs before transplantation, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities.
         08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
17-NOV-2000
17-NOV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel isolated human cytoskeletal element-related polypeptide useful f diagnosis/treatment of neoplastic disorders, disorders associated with neural transmission, chromosomal abnormalities, autoimmune disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
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2000US-0225267P
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2000US-02256868P
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2000US-023124178PP
2000US-0241785P
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RESULT 10
AAU51226
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copolypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO syndrome (synovitis, acne, pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central nervous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The colypeptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to downregulate expression and activity of P. acnes polypeptides and ctivity of P. acnes polypeptides and colypeptides and determining P. acnes presence, for example, by carpine linked immunosorbent assay (BLISA). Note: The sequence data for this patent did not form part of the printed specification, but was cobarined in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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02-JUN-2000; 2000US-0208841P.
07-JUL-2000; 2000US-0216747P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Propionibacterium acnes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        dermatological; osteopathic; neuroprotectant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        uveitis; endophthalmitis; bone; joint; central nervous system; E inflammatory lesion; acne vulgaris; enzyme linked immunosorbent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SAPHO syndrome; synovitis; acne;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Propionibacterium acnes immunogenic protein #12122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-FEB-2002
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                                                                                                                                                                                                                                                                                                       Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic polypeptides. The proteins and their associated DNA sequences are used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; AAS59551.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      L'maisonneuve
                                                                                                                                                                                                                                                                                                                                                         Example 1; SEQ ID NO 12421; 1069pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                          Propionibacterium acnes vaccinating against and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CORI-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IGDIGD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard; protein;
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                                                                                                                                                                                                                                                                                                                                                                                             acne vulgaris.
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e J, Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
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                                                                                                                                                                                                                                                                                                                                                                                                            polypeptides and nucleic acids useful for diagnosing infections, especially useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mitcham JL, Wang S., Jen S, Carter D;

    Mismatches

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Pred. No. 2.6e+02;
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RESULT 11
ABM47745
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CC encoding a Propionibacterium across protein. The invention also relates to CC polypeptides encoded by the polynucleotides (ABM35624-ABM64336) and to CC immunogenic fragments of P. across polypeptides. The invention additionally encompasses expression vectors and host cells comprising a cc polynucleotide of the invention, antibodies against polypeptides of the invention; a cc method for stimulating an immune response specific for a P. across CC polypeptide and an isolated T cell population comprising P. across CC polynucleotides, antibodies, fusion proteins, T cell populations, or CC polynucleotides, antibodies, fusion proteins, T cell populations, or CC polynucleotides, antibodies, fusion proteins, T cell populations, or CC patient; and a method for inhibiting the development of P. across in a CC patient; and a method for inhibiting the development of P. across in a CC patient; and a method for inhibiting the development of P. across the proteins, T cell populations or antigen-presenting cells that express the polypeptides, antibodies, fusion CC patient. The P. across polypeptides, polynucleotides, antibodies, fusion CC patient. The polynuclations or antigen-presenting cells that express the proteins, or for stimulations or antigen-presenting cells that express the polypeptides are useful for diagnosting, preventing or treating across the polypeptides are useful for across in a cc polypeptide are properly properly properly proteins. The polynucleotides can also be used as probes or primers for nucleic acid hybridisation. The vaccine composition is useful for the stimulation of an immune response against P. across, or for treating acros, and the kit is useful for performing a diagnostic assay. The present
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Best Local
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Zhang Y,
Barth B,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
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N-PSDB; ACF64480.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Acne vulgaris; antiseborrhoeic; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Propionibacterium acnes predicted ORF-encoded polypeptide #12421.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; SEQ ID NO 12421; 1481pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to an isolated polynucleotide (ACF64435-ACF64733)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mmunostimulant; immune
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49 VGDIGD 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     m JL, on. Jen s, Y, Wang S, Jen s, B, Vallieve-Douglass J;
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Jen S,
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Lodes MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 31; DB 4;
Pred. No. 3e+02;
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Benson DR,
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Jones R, Carter D;
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RESULT 12
ADUI8521
ID ABUI8521
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Best Local Similarity
Matches 5; Conserv
    31-JAN-2000
04-FEB-2000
24-FEB-2000
24-FEB-2000
16-MAR-2000
17-MAR-2000
17-MAR-2000
19-MAY-2000
07-JUN-2000
07-JUN-2000
07-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cytoskeletal element-related protein; human; mouse; rabbit; goat; horse; cat; dog; chicken; sheep; immunosuppressive; antiarthritic; vasotropic; antirheumatic; antiproliferative; cytostatic; cardiant; neuroprotective; cerebroprotective; nootropic; antibacterial; virucide; fungicide; cancer; ophthalmological; vulnerary; gene therapy; autoimmune disease; neoplasm; hyperproliferative disorder; brasst; liver; cardiovascular disorder; provous system disorder; bacterial infection; fungal infection; viral infection; ocular disorder; endocrine disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence represents a polypeptide predicted to be encoded by an (reading frame) contained within the P. across polypucleotides of trevention. Note: The sequence data for this patent did not form the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fungal infection; viral infection; ocular disorder; enc
gastrointestinal disorder; respiratory
wound healing; skin aging; organ transplantation; food
tissue regeneration; anti-infertility; food additive.
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    2000US-011965P.
2000US-0184664P.
2000US-0184659P.
2000US-0184659P.
2000US-0198123P.
2000US-0199875P.
2000US-0209467P.
2000US-021486P.
2000US-0216647P.
2000US-02174967P.
2000US-02174967P.
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2000US-02174967P.
2000US-02174967P.
2000US-02174967P.
2000US-02174967P.
2000US-02252164P.
2000US-0225213P.
2000US-0225214P.
2000US-0225214P.
2000US-0225266P.
2000US-0225266P.
2000US-0225270P.
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Pred. No. 3e+0
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 6;
3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                preservative;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disorder;
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t directly
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    14-AUG-2000
14-AUG-2000
11-AUG-2000
12-AUG-2000
22-AUG-2000
23-AUG-2000
23-AUG-2000
23-AUG-2000
23-AUG-2000
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2000US-0227182P

2000US-0229343P

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2000US-0231243P

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2000US-0244180P

2000US-0244178P

2000US-024647P

2000US-024652P

2000US-024652P
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01-DEC-2000;
01-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
06-DEC-2000;
08-DEC-2000;
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08-DEC-2000;
08-DEC-2000;
as glomerulonephritis and respiratory disorders such as glomerulonephritis and respiratory disorders such as asthma. The polypeptides can also be used to aid wound healing, to prevent skin aging due to sumburn, to maintain organs before transplantation, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published pct sequences.
                                                                                                                                                                                                               diseases such as rheumatoid arthritis, hyperproliferative disorders such as neoplasms of the breast or liver, cardiovascular disorders such as cardiac arrest, cerebrovascular disorders such as cerebral ischaemia, nervous system disorders such as Alzheimer's disease, infections caused by bacteria, viruses and fungi, ocular disorders such as corneal infection, endocrine disorders such as premature labour and infertility, gastrointestinal disorders such as Crohn's disease, renal disorders such as crops such as control disorders such as crops and disorders such as crops as the control of 
                                                                                                                                                                                                                                                                                                                                                                                             sequences AAU18508-AAU18551 represent the cytoskeletal element-related polypeptides of the invention. Cytoskeletal polypeptides and their associated polynucleotides are useful in the diagnosis, treatment and prevention of various types of disorders in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. A pathological condition can be determined by determining the presence or absence of a mutation in a cytoskeletal polynucleotide. The treatable disorders include autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated human cytoskeletal element-related polypeptide useful for diagnosis/treatment of neoplastic disorders, disorders associated with neural transmission, chromosomal abnormalities, autoimmune disorders.
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2000US-0251868P
2000US-0251869P
2000US-0251989P
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for

Query Match

96.98;

Score 31;

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19-JUL-2000;
01-SEP-2000;
15-SEP-2000;
20-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; cytokine; cell proliferation; cell differentiation; gene th vaccine; peptide therapy; stem cell growth factor; haematopolesis; tissue growth factor; immunomodulatory; cancer; leukaemia;
                       The invention relates to polynucleotides (AAKS1456-AAKS3435) and the encoded polypeptides (AAM78323-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polymerides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, immunomodulatory activity and activity and may be useful in the diagnosis and/or treatment of cancer, leukaemia, nervous system disorders, arthritis and inflammation. Note: Records for SEQ ID NO 2110 (AAKS2581), 2111 (AAKS2582) and 3666 (AAM80020) are omitted as the relevant pages from the sequence listing were missing at the time of publication
                                                                                                                                                                                                                                                                                                                                Tang
Ma Y,
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                                                                                                                                                                                                           Claim 20; Page 3461-3462; 6221pp;
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                                                                                                                                                                                                                                   Nucleic acids encoding polypeptides with cytokine-like activities, useful in diagnosis and gene therapy.
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Sequence 166 AA;
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5; Conser
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; 2000US-00560875.

; 2000US-00598075.

; 2000US-00620325.

; 2000US-00634936.

; 2000US-00633251.

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; 2000US-00728422.
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Pred. No.
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J J, Zhang Goodrich R
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3.1e+02;
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Best Local
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Matches
                                                                                                                                                                The present sequence is the protein sequence of mouse cofilin. This is encoded by a gene that has been identified as being up-regulated in 2 models of chronic pain, i.e. streptozocin-induced diabetes and chronic constrictive injury to a nerve leading to the spine. The expression products of such genes can be used to screen libraries for compounds and peptide agonists and antagonists of gene product activity that may be useful in the treatment or prevention of chronic pain, and in the development of diagnostic tools for the identification and
                                                                                                                                                                                                                                                                      Use of isolated gene or nucleic acid sequence, recombinant vector, host cell, non-human animal, polypeptide encoded by the nucleic acid sequence antibody, for screening of compounds for the treatment of pain, or fi
                                                                                                                                                                                                                                                                                                                                                                     27-JUL-2001; 2001GB-00018354.
07-FEB-2002; 2002GB-00002892.
         ADB75248;
                                                                                                                                        Sequence 166 AA;
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                          ADB75248 standard;
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1; Mismatches 0
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Pred. No. 3.3e+02;
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Best Local Similarity
Matches 5; Conserv
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22-AUG-2001; 2001US-0314356P.
25-SEP-2001; 2001US-0325020P.
12-DEC-2001; 2001US-0341746P.
05-MAR-2002; 2002US-0362158P.
                                                                                                                                                                                                             Sequence 166 AA;
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                                                            06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                      cancer.
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83.3%;
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assessing whether a patient is afflicted with prostate cancer. The method of the invention involves assessing whether a patient is afflicted with prostate cancer by comparing the level of expression of a marker in a patient sample and the normal level of expression of the marker in a control non-prostate cancer sample, where a significant increase in the level of expression of the marker in the patient sample and the normal level of expression of the marker in the patient sample and the normal level indicates that the patient is afflicted with prostate cancer. Nucleic acids of the invention are useful for diagnosing or treating prostate cancer, and may be useful in gene therapy. Sequences given in ADB75177-ADB75631 represent marker cDNA and proteins. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at
mitochondrial; human; screening assay; diabetes mellitus; Huntington's disease; osteoarthritis; Leber's hereditary optic neuropathy; LHON; mitochondrial encephalopathy lactic acidosis and stroke;
                                                                                                                                                                                   Human heat mitochondrial protein as a therapeutic target SeqID1910.
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                                                                                                                                                                                                                                                                                                                                                                                                             ADJ70104 standard; protein; 166 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wonsey AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.
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Pred. No. 3.3e+02;
1; Mismatches 0
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M, Glatt K, Zhao
   acidosis and stroke; MELAS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 166;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                or
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Σ,
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Anderson D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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RESULT 17
ADI24536
ID ADI24
XX
AC ADI24
XX
DT 15-AF
DT 15-AF
XX
Chkl
KW Chkl
KW gene
XX
OS Homo
XX
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                                                                                                                                                                                                                                                                                                                                                                                                    CC can be used in the treatment of various disease associated with altered mitochondrial function. Specifically, it refers to a method for cuseful for drug screening assays, as well as therapeutic targets. The can be used in the treatment of various diseases associated with the can be used in the treatment of various diseases associated with altered cc can be used in the treatment of various diseases associated with altered cc osteoarthritis, Leber's hereditary optic neuropathy (LHON), mitochondrial cc encephalopathy lactic acidosis and stroke (MELAS), myoclonic epilepsy cc ragged red fibre syndrome (MERRF) or cancer. Accordingly, these cc compositions have neuroprotective, nootropic, antidiabetic, antiarthritic, osteopathic, ophthalmological and cc cytostatic activities. This polypeptide sequence is a human heart cc mitochondrial protein of the invention.
                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-APR-2002;
17-JUN-2002;
20-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                myoclonic epilepsy ragged red fibre syndrome; MERRF; cancer; neuroprotective; nootropic; antidiabetic; anticonvulsant; antiarthritic; osteopathic; ophthalmological; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying a mitochondrial target for drug screening assays and for treating diseases associated with altered mitochondrial function, comprises detecting a modified polypeptide in a sample and correlating
                                                                                                                                                                                                                                                                                                                                                                            Sequence 166 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to novel mitochondrial targets that can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 1910; 180pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-845369/78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ghosh SS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-APR-2003; 2003WO-US010870.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MITO-) MITOKOR. (BUCK-) BUCK INS
                                                                                         Human modifier of Chk1 (MCHK) protein SEQ ID NO:86
                 Homo sapiens
                                                                                                                     15-APR-2004
                                                                                                                                                                               ADI24536 standard; protein; 166 AA.
                                                                                                                                                                                                                                                                                                               Local Similarity hes 5; Conserv
                                          pathway modulating agent; modifier of Chk1; MCHK; therapy; cancer; human.
                                                                                                                                                                                                                                                       57 VGDIGD 62
                                                                                                                                                                                                                                                                                   1 ignign 6
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                                                                                                                                                                                                                                                                                                                   Conservative
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2002US-0389987P.
2002US-0412418P.
                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                 96.9%;
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Pred.
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                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                  31;
No.
                                                                                                                                                                                                                                                                                                                                  3.3e+02;
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                                                             cytostatic;
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RESULT 18
AAU52312
ID AAU52
XX AAU52
XX AAU52
XX 27-FE
XX SAPHO
KW UVE:IT
KW infla
XX UVE:IT
KW infla
XX Propi
XX ORTHO

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Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a method for identifying a candidate Chk1 pathway modulating agent. The method comprises: (a) providing an assay system comprising a modifier of Chk1 (MCHK) polypeptide or nucleic acid; (b) contacting the system with a test agent, where the system provides a reference activity except in the presence of the test agent; and (c) detecting a test agent-biased activity, and a difference between the test agent-biased activity and the reference activity. Also described: (1) a method for modulating Chk1 pathway of a cell; (2) a method for modulating Chk1 pathway in a mammalian cell; and (3) a method for diagnosing a disease in a pathent. A MCHX sequence has cytostatic activity, and can be used in gene therapy. The method is useful for identifying a candidate chk1 pathway modulating agent for preparing a composition for diagnosing or treating e.g., cancer. The present sequence represents a human MCHX protein, which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying a candidate Chkl pathway modulating agent for treating e.g., cancer, comprises contacting an assay system comprising a MCHK polypeptide or nucleic acid with a test agent and detecting a test agent-biased activity.
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N-PSDB; ADI24486.
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16-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAU52312 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EXEL-)
                                                                                                                                                                                                                                                                                                                                                   Propionibacterium acnes immunogenic protein #13208
                                                                                                                                                                                                                                                                                                                                                                                                              27-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAU52312;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               166 AA;
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2002US-0410986P
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Amundsen CD,
                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              62
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Jin
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Adamkewicz
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ა.3e+02;
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Platt
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att DM;
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SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay;

osteopathic; acnes

20-APR-2001; 2001WO-US012865.

01-NOV-2001

Propionibacterium dermatological;

WO200181581-A2

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC polypeptides. The proteins and their associated DNA sequences are used in CC the treatment, prevention and diagnosis of medical conditions caused by CP. acnes. The disorders include SAPHO syndrome (synovitis, acne, compared by CC pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. CC pervous system, however it is particularly involved in the inflammatory CC lesions associated with acne vulgaris. A method for detecting the CC presence or absence of P. acnes in a patient comprises contacting a CC sample with a binding agent that binds to the proteins of the invention CC and determining the amount of bound protein in the sample. The CC polypeptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to CC downregulate expression and activity of P. acnes polypeptides and CC therefore treat P. acnes infections. The antibodies may also be used as diagnostic agents for determining P. acnes presence, for example, by C enzyme linked inmunosorbent assay (ELISA). Note: The sequence data for this patent did not form part of the printed specification, but was CC obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Propionibacterium acnes polypeptides and nucleic acids useful for vaccinating against and diagnosing infections, especially useful for treating acne vulgaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-APR-2000; 2000US-0199047P.
02-JUN-2000; 2000US-0208841P.
07-JUL-2000; 2000US-0216747P.
15-OCT-2001; 2001US-00978825
                               11-OCT-2002; 2002WO-US032727
                                                                                                WO2003033515-A1
                                                                                                                              Propionibacterium acnes.
                                                                                                                                                                         Acne vulgaris; antiseborrhoeic; dermatological; antibacterial;
                                                                                                                                                                                                          Propionibacterium acnes predicted ORF-encoded polypeptide #13507
                                                                                                                                                                                                                                             20-OCT-2003
                                                                                                                                                                                                                                                                            ABM48831;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; SEQ ID NO 13507; 1069pp; English
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                                                                                                                                                             immunostimulant;
                                                                                                                                                                                                                                                                                                         ABM48831 standard; protein; 173 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                       107
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                                                                                                                                                             immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   96.9%;
                                                                                                                                                             response;
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, Jen S, Carter D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 31; DB 4; : Pred. No. 3.5e+02; 1; Mismatches 0
                                                                                                                                                               vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 173;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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RESULT 20
ADQ10039
ID ADQ10
XX ADQ10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                         variable major protein; VMP; VMP-like sequence; VLS; Lyme disease;
                                                                                                                                                                                                                                                                                                  B_afzelii ORF 13 protein product of the vls locus silent cassette Seq 83.
                                                                                                                                                                                                                                                                                                                                                                        07-OCT-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                           ADQ10039;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADQ10039 standard; protein; 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 173 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
                                                                                                  Borrelia afzelii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; SEQ ID NO 13507; 1481pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mitcham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                             lapsing fever; antibacterial; gene therapy; antigenic variation system; pathogenic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2003-381789/36.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
5; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JĿ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wang S, Jen S, I
Vallieve-Douglass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Skeiky YAW,
ing S, Jen S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   96.9%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Persing DH,
Lodes MJ,
188 J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 31; DB 6;
Pred. No. 3.5e+02;
1; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bhatia A,
Benson DR,
                                                                                                                                                                                                immunoprophylaxis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Maisonneuve JL;
Jones R, Carter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     æ
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ARBSULT 21
AAU18546
ID AAU18546
AC AAU18
XX AU18
AC AAU18
XX C1-NC
DT 21-NC
DT 21-NC
CYTOE
KW CYTOE
KW CAT:
KW CHING
KW CETEL
KW ANT:
KW ANT:
KW ANT:
KW Hypes
KW CETEL
KW GETEL
KW GETEL
KW GETEL
KW GETEL
KW GUNN
KW HOMO
OS HOMO
PN WO20
PN WO20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention relates to a novel nucleic acid molecules isolated from CC pathogenic Borrelia bacteria that each encode a variable major protein CC (VMP)-like sequence (VLS) peptide. Specifically, it refers to recombinant CC VLS genes and immunogenic compositions derived thereof, from Borrelia CC garinii (B. garinii) and Borrelia afzelii (B. afzelii). The present CC invention describes the use of antibodies against the expressed VLS CC proteins where immunologic binding is indicative of a Borrelia infection, CC in particular infections that include Lyme disease, relapsing fever or CC related diseases in humans and animals. Accordingly, these antibacterial CC compositions can be used for gene therapy, immunoprophylaxis, diagnosis CC or treatment of such diseases, which due to the VMP antigenic variation CC system often evade the immune system response despite the occurrence of host antibody and cellular responses. This polypeptide is a protein CC encoded by a Borrelia afzelii strain ACAI open reading frame DNA sequence CC from the VIs locus silent cassette of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                         cytoskeletal element-related protein; human; mouse; rabbit; goat; horse; cat; dog; chicken; sheep; immunosuppressive; antiarthritic; vasotropic; antirheumatic; antiproliferative; cytostatic; cardiant; neuroprotective; cerebroprotective; nootropic; antibacterial; virucide; fungicide; cancer; ophthalmological; vulnerary; gene therapy; autoimmune disease; neoplasm; hyperproliferative disorder; breast; liver; cardiovascular disorder; hyperprosecular disorder; breast; liver; cardiovascular disorder; fungal infection; fungal infection; viral infection; ocular disorder; endocrine disorder; gastrointestinal disorder; renal disorder; respiratory disorder; wound healing; skin aging; organ transplantation; food preservative; tissue regeneration; anti-infertility; food additive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid molecules encoding Variable Major Protein-like polypeptides of Borrelia, useful for diagnosing, preventing or treating Borrelia infections such as Lyme disease or relapsing fever.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 22; SEQ ID NO 83; 182pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Norris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-DEC-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
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                                                                                                                                                                                                                                                                                                                                                                                          Human cytoskeletal element-related polypeptide #39.
                                                                                                                                                                                                                                                                                                                                                                                                                                    21-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAU18546;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAU18546 standard; protein; 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 201
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    02-AUG-2001
                                               WO200155168-A1
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DB; ADQ10038.
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5; Conserv
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IGDVGD 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           96.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 31; DB 8;
Pred. No. 4e+02;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 201;
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29-SEP-2000;
22-SEP-2000;
22-CCT-2000;
22-CCT-2000;
22-CCT-2000;
23-CCT-2000;
23-CCT-2000;
23-CCT-2000;
23-CCT-2000;
20-CCT-2000;
20-NOV-2000;
21-NOV-2000;
21-NO
WPI; 2001-476182/51.
N-PSDB; AAS29808.
                                                                                       Rosen CA,
                                                                                                                                                     HUMAN
                                                                                       Barash SC,
                                                                                                                                                                                                            2000US-0236369P.
2000US-0236802P.
2000US-0236802P.
2000US-0236802P.
2000US-0237039P.
2000US-023937P.
2000US-02499399.
2000US-0241786P.
2000US-0241809P.
2000US-0241809P.
2000US-0246474P.
2000US-0246474P.
2000US-0246477P.
2000US-0246477P.
2000US-0246477P.
2000US-0246478P.
2000US-0246523P.
2000US-0246610P.
2000US-0246611P.
2000US-0246611P.
2000US-0246611P.
2000US-0246611P.
2000US-024921P.
2000US-0251186.
2000US-02
                                                                                                                                                     GENOME
                                                                                                                                                     SCI INC
                                                                                              Ruben SM;
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Query Match
Best Local
                                                                                                                                                                                                                                                                                                                             polypeptides of the invention. Cytoskeletal polypeptides and their associated polymucleotides are useful in the diagnosis, treatment and prevention of various types of disorders in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. A pathological condition can be determined by determining the presence or absence of a mutation in a cytoskeletal polymucleotide. The treatable disorders include autoimmune diseases such as rheumatoid arthritis, hyperproliferative disorders such as neoplasms of the breast or liver, cardiovascular disorders such as cardiac arrest, cerebrovascular disorders such as cerebral ischaemia, nervous system disorders such as Alzheimer's disease, infections caused by bacteria, viruses and fungi, ocular disorders such as corneal infection, endocrine disorders such as premature labour and infertility, gastrointestinal disorders such as Crohn's disease, renal disorders such as gastrointestinal disorders such as Crohn's disease, renal disorders such as consecutions.
                                                                                      specification, at ftp.wipo.int
                                                                                                                   as glomerulonephritis and respiratory disorders such as asthma. The polypeptides can also be used to aid wound healing, to prevent skin aging due to subburn, to maintain organs before transplantation, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequences AAU18508-AAU18551 represent the cytoskeletal element-related polypeptides of the invention. Cytoskeletal polypeptides and their associated polynucleotides are useful in the diamnosis transmission.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel isolated human cytoskeletal element-related polypeptide useful for diagnosis/treatment of neoplastic disorders, disorders associated with
                                                                                         ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11; SEQ ID NO 93; 505pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         transmission,
96.9%;
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                            Length 205;
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RESULT 22
AAM79529
ID AAM79
XX AAM79
AC AAM79
AC AAM79
XX U6-NQ
CX Humar
XX Humar
XX Homo
XX Homo
XX Homo
XX Homo
XX Homo
XX 109-Al
XX 03-FE
PR 27-Al
PR 19-UI
PR 19-UI
PR 19-UI
PR 19-UI
PR 119-UI
PR 210-OS
PR 20-NG
XX PA
PR 20-NG
XX PA
PR 20-NG
XX PA
PR 20-NG
PR 30-NG
XX PA
PR 20-NG
PR 30-NG
XX PA
PR 30-N
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                                                                                                              03-FEB-2000; 2000US-00496914.

27-APR-2000; 2000US-00560875.

20-JUN-2000; 2000US-00598075.

19-JUL-2000; 2000US-00620325.

01-SEP-2000; 2000US-00620325.

15-SEP-2000; 2000US-00634936.

15-SEP-2000; 2000US-00693325.

20-CCT-2000; 2000US-00728422.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; cytokine; cell proliferation; cell differentiation; gene th vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAM79529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-FEB-2001; 2001WO-US004098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200157190-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity tes 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 IGDIGD 6
:|||||
96 VGDIGD 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein SEQ ID NO 3175.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             system disorder; arthritis; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; protein; 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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Pred. No. 4.1e+02;
1; Mismatches 0
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(HYSE-) HYSEQ

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RESULT 23
ADQ10053
ADQ10053
ADQ100
XX ADQ10
XX ADQ10
XX ADQ10
XX Varia
XX Varia
XX VAP
XX VAP
XX VAP
XX SBORY
XX WO20
PN W020
PN W220
PF 22-D
XX 22-D
XX ATA
PF NOT
PR NOT
XX NP
DR WPI
DR WPI
DR WPI
DR N-PI
XX NP
PT New
PT New
PT New
PT PO1
PS Cla
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Ma Y, .
e AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            encoded polypeptides (AAM78333-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polymucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, immunomodulatory activity and activin/inhibin activity and may be useful in the diagnosis and/or treatment of cancer, leukaemia, nervous system disporders, arthritis and inflammation. Note: Records for SEQ ID NO 2110 (AAK52581), 2111 (AAK52582) and 3666 (AAM80020) are omitted as the relevant pages from the sequence listing were missing at the time of publication
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-476283/51.
N-PSDB; AAK52662.
                                                                                                                                                             WPI; 200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Borrelia afzelii.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 variable major protein; VMP; VMP-like sequence; VLS; Lyme disease; relapsing fever; antibacterial; gene therapy; immunoprophylaxis; VMP antigenic variation system; pathogenic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B_afzelii strain ACAI vls 13
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                                                New isolated nucleic acid molecules encoding Variable Major polypeptides of Borrelia, useful for diagnosing, preventing Borrelia infections such as Lyme disease or relapsing fever.
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7, Zhao QA,
AJ, Yang Y,
                                                                                                                                                          2004-525783/50.
DB; ADQ10052.
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J, Zhang
Goodrich R
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odrich R;
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J, Ren 1
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Claim 22; SEQ ID NO

97; 182pp; English

The invention relates to a recombinant DNA construct comprising a promoter functional in plants operably linked to an oil-associated gene. The construct is useful for transgenic plant seed which has in its genom the construct, that is functional in the plant to transcribe the oil-

its genome he oil-

Disclosure;

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Matches
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                                                                                                                                                                                   Novel recombinant DNA construct comprising a promoter functional in plants operably linked to an oil-associated gene for producing transplant seed.
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(RAVA/)
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26-JUN-2002; 2002US-0391786P.
26-JUN-2002; 2002US-0392018P.
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RAVANELLO M.
SAVAGE T.
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Best Local (
                 The invention relates to a recombinant DNA construct comprising a promoter functional in a plant cell, where the promoter is positioned to provide for expression of a polymucleotide encoding a polympetide from a microbial source. The invention also relates to a transformed plant comprising the recombinant DNA construct and a method of producing a transformed plant having an improved property. The plant is a crop plant such as maize or soybean. The method of producing a transformed plant with the recombinant DNA construct and growing the transformed plant with the polymucleotide or polypeptide is useful for improving plant properties. The recombinant DNA construct is useful for improving plant properties.
                                                                                                                                                                                                                                                                                                                                                                                                                         New recombinant DNA construct comprising a promoter positioned to provide for expression of a polynucleotide encoding a polypeptide from a microbial source, useful for producing plants with improved properties.
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(HINK/)
(SLAT/)
(CHEN/)
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HINKLE G
SLATER S
CHEN X.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phosphorus; photosynthesis; lignin; galactomannan;
polypeptide.
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Pred. No. 6.9e
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6.9e+02;
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  drought tolerance
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            δ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tolerance to herbicides, extreme osmotic conditions, pathogens or pests, increased resistance to plant disease, better growth rate by modification of the cell cycle pathway with plant growth regulators, increased rate of homologous recombination, modified seed oil or protein yield and/or content, improved yield by modification of carbohydrate, nitrogen or phosphorus use and/or uptake, by modification of photosynthesis or by providing improved plant growth and development under at least one stress condition, improved lignin production or improved galactomannan production. This sequence represents a bacterial polypeptide used in the scope of the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Recombinant DNA construct; transformed plant; improved plant property; cold tolerance; heat tolerance; drought tolerance; herbicide; osmosis; pathogen tolerance; pest tolerance; plant disease resistance; pathogen tolerance; plant growth regulator; cell cycle pathway modification; plant growth regulator; homologous recombination; seed oil yield; protein yield; carbohydrate;
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(SLAT/)
(CHEN/)
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SLATER S
CHEN X.
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The invention relates to a recombinant DNA construct comprising a promoter functional in a plant cell, where the promoter is positio provide for expression of a polynucleotide encoding a polynucleotid

comprising a ter is positioned to

plant

from þ New recombinant DNA construct comprising a promoter positioned to provide for expression of a polynucleotide encoding a polypeptide from a microbial source, useful for producing plants with improved properties.

SEQ ID NO 11876; 122pp; English.

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RESULT 27
AAB22942
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Matches 5
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This sequence represents mouse GANP protein. The invention relates to the mosue and human GANP proteins (AAB22942, AAB22943) which are 210 kD in size, have kinase activity and participate in signal transduction during abnormal B cell differentiation. The invention also encompasses nucleotides encoding GANP proteins, variant GANP proteins, and antibodies
                                                                                                                                                                                                                                                  GANP proteins participating in signal conversion of abnormal B cell differentiation in autoimmune state and having kinase activity, use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-AUG-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                          Sakaguchi N, Kuwahara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-FEB-1999;
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                                                                                                                                                                        Claim 1; Page 58-67; 91pp; Japanese.
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83.3%;
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RESULT 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     against GANP proteins. GANP proteins may be used mechanisms involved in autoimmune disorders
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                                                                                                                                                                                                                                                                                                                                         WPI; 2001-313371/33.
N-PSDB; AAH25842.
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                                                                                                                                                                                                               Sequence 1971 AA;
                                                                                                                                                                                                                                 AAB98973. This protein can be used for controlling diffe activation of germ centre B cells. The present sequence in the exemplification of the invention
                                                                                                                                                                                                                                                          The present invention relates to the murine SHD1 protein shown in AAB98973. This protein can be used for controlling differentiation and
                                                                                                                                                                                                                                                                                      Claim 2; Page 8-13; 16pp;
                                                                                                                                                                                                                                                                                                          SHD1 protein useful for controlling the differentiation germ center B cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse; SHD1;
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                                                                                                                                                                                                                                                                                                                                                                                           17-SEP-1999;
Mouse; CALLP; autoimmune disease.
                    Murine CALLP related protein
                                        20-AUG-2001
                                                            AAB98957;
                                                                               AAB98957 standard;
                                                                                                                                                                                                                                                                                                                                                                       (SUME ) SUMITOMO ELECTRIC IND
                                                                                                                                                                                 Local Similarity
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                                                                                                                                  1388 VGDIGD 1393
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5; Conserv
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                                        (first entry)
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                                                                              protein;
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83.3%;
                                                                                                                                                                                   96.9%;
                                                                                                                                                                                                                                                                                        Japanese.
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                                                                                                                                                                                   Score 31;
Pred. No.
                                                                                 1971 AA
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Pred. No. 4.2e+03;
1; Mismatches 0
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                                                                                                                                                                           Mismatches
                                                                                                                                                                                     4.2e+03;
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RESULT 30
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Best Local Similarity
Matches 5; Conservi
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      Transgenic mammal transformed with germinal center associated nuclear protein (GANP) gene for production of high-affinity antibodies as diagnostic reagents and disease therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides a murine protein, designated CALPP and shown in AAB9896. This can be used for the treatment or detection of autoimmune diseases. The present sequence is a protein used in the exemplification of the
                                                                                                                                                                                                      Mus
                                                                                                                                                                                                                         germinal center associated nuclear protein; GANP; antibody; virus infection; human immunodeficiency virus; hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1971 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New CALPP protein the use of which is detection of autoimmune diseases.
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                                                                              Sakaguchi N;
                                                                                                                     07-NOV-2002; 2002WO-JP011598
                                                                                                                                        07-NOV-2003; 2003WO-JP014221
                                                                                                                                                                                                                                            virucide; viral antigen inhibitor; transgenic;
                                                                                                                                                                                                                                                                Mouse germinal center associated nuclear protein
                                                                                                                                                                                                                                                                                      12-AUG-2004
                                                                                                                                                                                                                                                                                                                            ADO31144 standard; protein; 1971 AA
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                                                                                                  (-DWMI)
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                                                          2004-411378/38
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                                                                                                                                                                                                                                                                                                                                                                                                  1 IGDIGD
                                                                                                  IMMUNOKICK INC
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83.3%;
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4.2e+03;
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KW Infla
KW O200
XX Propi
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XX O1-NC
XX O200
PR 02-JI
PR 02-JI
PR 07-JI
PR 07-JI
PR 07-JI
PR N-PSI
XX ICOR:
XX WPI;
PI L'ma:
XX WPI;
PI Skeil
PI Kreai
XX Propi
XX SAPHC
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XX O200
PF 20-AI
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Best Local
                        Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogeni polypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO syndrome (synovitis, acne, pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central nervous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to transgenic non-human animals and their offspring which are transformed with germinal center associated nuclear protein (GANP) gene. The GANP gene, encoded protein and transgenic animals express GANP can be used for the production of high-affinity antibodies to viral antigens for treatment and prevention of infection by viruses such as human immunodeficiency virus and hepatitis C virus. This sequence
                                                                                                                                                                                                                                                                                                                                                                                                                      Propionibacterium acnes polypeptides and nucleic acids useful for vaccinating against and diagnosing infections, especially useful for treating acne vulgaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-616774/71.
N-PSDB; AAS59663.
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02-JUN-2000; 2000US-0208841P.
07-JUL-2000; 2000US-0216747P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; ELISA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                             Example 1; SEQ ID NO 26477; 1069pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-APR-2001; 2001WO-US012865
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay;
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Zhang Y,
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, Jen S, Carter
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Pred. No. 4.2e
1; Mismatches
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4.2e+03;
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are used in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mitcham JL, Ske
Zhang Y, Wang S
Barth B, Vallie
The invention relates to an isolated polynucleotide (ACF64435-ACF64733) encoding a Propionibacterium acnes protein. The invention also relates to polypeptides encoded by the polynucleotides (ABM35624-ABM64536) and to immunogenic fragments of P. acnes polypeptides. The invention and to immunogenic fragments of P. acnes polypeptides. The invention additionally encompasses expression vectors and host cells comprising a polynucleotide of the invention; antibodies against polypeptides of the invention; a method for stimulating an immune response specific for a P. acnes polypeptide and an isolated T cell population comprising T cells prepared via this method; a vaccine composition (comprising P. acnes polypeptides, antibodies, fusion proteins, T cell populations, or antigen-presenting cells that express the polypeptide); a method and kit for detecting or determining the presence or absence of P. acnes in a patient; and a method for inhibiting the development of P. acnes in a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New Propionibacterium acnes polypeptides and polynucleotides encoding the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; SEQ ID NO 26477; 1481pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CORI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ypeptide, useful for diagnosing, preventing for stimulating an immune response specific
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACF64592.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LGDIGD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vallieve-Douglass J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Skeiky YAW,
ng S, Jen S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93.8%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        response;
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Lodes MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 30; DB of Pred. No. 1.8e.
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Benson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bhatia A,
Benson DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  or treating acne vulgaris, for a P. acnes protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Maisonneuve JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ₽,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Carter D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogen. polypeptides. The proteins and their associated DNA sequences are used the treatment, prevention and diagnosis of medical conditions caused by p. acnes. The disorders include SAPHO syndrome (synovitis, acne, pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central P. acnes is also involved in infections of bone, joints and the central P. acnes is also involved in infections of bone, joints and the central P. acnes is also involved in infections of bone, joints and the contral P. acnes is also involved in infections of bone, joints and the contral p. acnes is also involved in infections of bone, joints and the contral p. acnes is also involved in infections of bone, joints and the contral p. acnes is also involved in infections of bone.

Example 1; SEQ ID NO 20489;

1069pp;

English.

immunogenic are used in

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AAU59294
ID AAU
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RESULT 33
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Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           patient. The P. acnes polypeptides, polynucleotides, antibodies, fusion proteins, T cell populations or antigen-presenting cells that express the polypeptides are useful for diagnosing, preventing cells that express the polypeptides are useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein. The polynucleotides can also be used as probes or primers for nucleic acid hybridisation. The vaccine composition is useful for the stimulation of an immune response against P. acnes, or for treating acne, and the kit is useful for performing a diagnostic assay. The present sequence represents a polypeptide predicted to be encoded by an ORF (open reading frame) contained within the P. acnes polynucleotides of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly form.
                                                                                                                                                                                                                                                                                                                  21-APR-2000; 2000US-0199047P.
02-JUN-2000; 2000US-0208841P.
07-JUL-2000; 2000US-0216747P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SAPHO syndrome; synovitis; acne; uveitis; endophthalmitis; bone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 58
                                                                                                                                                       Propionibacterium acnes vaccinating against and
                                                                                                                                                                                                  N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                             01-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200181581-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Propionibacterium acnes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                dermatological; osteopathic; neuroprotectant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Propionibacterium acnes immunogenic protein #20190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAU59294;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU59294 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  from WIPO at
                                                                                                                                         vaccinating against and
treating acne vulgaris.
                                                                                                                                                                                                                                            L'maisonneuve
                                                                                                                                                                                                                                                             Skeiky YAW,
                                                                                                                                                                                                                                                                                                                                                                              20-APR-2001; 2001WO-US012865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay;
                                                                                                                                                                                                                                                                                       (CORI-) CORIXA CORP.
                                                                                                                                                                                                   2001-616774/71.
DB; AAS59601.
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5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry
                                                                                                                                                                                                                                                             Persing
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                                                                                                                                                                                                                                              Zhang
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83.3%;
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                                                                                                                                                                                                                                               ĸ
                                                                                                                                                       polypeptides and nucleic acids useful for diagnosing infections, especially useful:
                                                                                                                                                                                                                                              Mitcham JL, Wang SS,
, Jen S, Carter D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 30; DB 6; Le
Pred. No. 1.8e+02;
1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pustulosis; hypertosis; osteomyelitis; joint; central nervous system; ELISA;
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                                                                                                                                                                                                                                                              Bhatia
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RESULT 34
ABM55813
ID ABM55813
AC ABM55
AC ABM55
AC ABM55
AC ACM5
AC Propi
XX Acme
KW Acme
KW Acme
KW Acme
KW Acme
CC PR 11-00
XX Propi
XX ACM5
PF 11-00
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XX ACM5
PF 11-00
XX Propi
XX Acm6
PF 11-00
XX Propi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mitcham JL,
Zhang Y, Wa
Barth B, Va
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The polypeptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to downregulate expression and activity of P. acnes polypeptides and therefore treat P. acnes infections. The antibodies may also be used as diagnostic agents for determining P. acnes presence, for example, by enzyme linked immunosorbent assay (ELISA). Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at
encoding a Propionibacterium acnes protein. The invention also relates to polypeptides encoded by the polynucleotides (ABM35624-ABM64336) and to immunogenic fragments of P. acnes polypeptides. The invention additionally encompasses expression vectors and host cells comprising a polynucleotide of the invention; antibodies against polypeptides of the invention; fusion proteins comprising a polypeptide of the invention; a method for stimulating an immune response specific for a P. acnes polypeptide and an isolated T cell population comprising P. acnes polypeptides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Acne vulgaris; antiseborrhoeic; dermatological; antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-OCT-2003
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                                                                                                                                                                                                                                                                                                                           The invention relates to an isolated polynucleotide (ACF64435-ACF64733)
                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; SEQ ID NO 20489; 1481pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunostimulant; immune response; vaccine.
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Vallieve-Douglass
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Lodes MJ,
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Pred. No. 2.4e+02;
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Benson DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Benson
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Jones R, Carter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cc polymucleotides, antibodies, fusion proteins, T cell populations, or cantigen-presenting cells that express the polypeptide); a method and kit cfor detecting or detecrining the presence or absence of p. acnes in a cc patient; and a method for inhibiting the development of p. acnes in a cpatient. The p. acnes polypeptides, polymucleotides, antibodies, fusion cc proteins, T cell populations or antigen-presenting cells that express the polypeptides are useful for diagnosing, preventing or treating acne correction. The polymucleotides can also be used as probes or primers for mucleic acid hybridisation. The vaccine composition is useful for the stimulation of an immune response specific for a p. acnes correction. The polymucleotides can also be used as probes or primers for the stimulation of an immune response against p. acnes, or for treating acne, cand the kit is useful for performing a diagnostic assay. The present canding frame) contained within the p. acnes polymucleotides of the creation. Note: The sequence data for this patient did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                    04-MAR-2002; 2002US-0361742P.
14-MAY-2002; 2002US-0380423P.
16-SEP-2002; 2002US-0410974P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hyperthermophile; protein stability enhancement; protein activity enhancement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADM27005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADM27005 standard; protein; 87 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 78 AA;
                                                                                                                                                                                                                         N-PSDB;
                                                                                                                                                                                                                           WPI; 2003-748383/70.
N-PSDB; ADM27081.
                                                                                                                                                                                                                                                                                Slesarev AI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-MAR-2003; 2003WO-US006664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Methanopyrus kandleri.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hyperthermophile Methanopyrus kandleri protein #1611.
                                                                                                                                                                                                                                                                                                                   (FIDE-) FIDELITY SYSTEMS (MALY/) MALYKH A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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71 VGDVGD
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                                                                                                                                                                                                                                                                                Pavlov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  93.8%;
                                                                                                                                                                                                                                                                                P
                                                                                                                                                                                                                                                                                                                                    INC
                                                                                                                                                                                                                                                                                Pavlova
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2; Mismatches
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Pred.
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                                                                                                                                                                                                                                                                                Kozyavkin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB
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The invention comprises the amino acid sequence of proteins from the hyperthermophile Methanopyrus kandleri, the invention also comprises the complete genome from Methanopyrus kandleri. The Methanopyrus kandleri proteins of the invention are useful for enhancing the stability and/or activity of other proteins. The Methanopyrus kandleri genome is useful in

New isolated nucleic acids kandleri proteins, and the as diagnostic agents.

encoding any of about 1700 Methanopyrus encoded proteins, useful as a medicamen

useful as a medicaments or

Claim 31; SEQ ID NO 1611; 1023pp; English

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The present invention describes a nucleic acid molecule (I) selected from CC DP119, DP444, DP810, DP685, WE474, DP160, RA977 and RA770, or its encoded CC polypeptide, or a fragment or variant of the nucleic acid or polypeptide, (CC or an effector/modulator of the nucleic acid or polypeptide. (I) is used CC in manufacturing a pharmaceutical agent, in monitoring or controlling the CC function of a gene or gene product influenced or modified by a DP119, CC DP444, DP810, DP685, WE474, DP160, RA977 or RA770 polypeptide. (I) has CC in gene therapy. (I) is useful for diagnostic or therapeutic CC applications, or for manufacturing an agent for diagnosts, monitoring, CC prevention or treatment of pancreatic disorders, such as diabetes and correct consequence of the correct of iseases. It may also be used for detecting and/or verifying, for the creatment, alleviation and/or prevention, of a pancreatic dysfunction CC (e.g. diabetes, hyperglycaemia or impaired glucose tolerance), and cc related disorders including obesity, and neurodegenerative disorders, and other contents, in cells, cell masses, organs and/or subjects. It is also used conters, in cells, cell masses, organs and/or subjects. It is also used contents and of the contents and or prevention of the cells, cell masses, organs and/or subjects. It is also used contents and or function of the contents and or function of the contents and or function of the contents and or promoting the differentiation and/or function of the contents and or promoting the differentiation and/or function of the contents and or promoting the differentiation and/or function of the contents and or promoting the differentiation and/or function of the contents and or promoting the differentiation and/or function of the contents and or promoting the differentiation and/or function of the contents and the con
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Use of DP119, DP444, DP810, DP685, WB474, DP160, RA977 or RA770 nucleic acid molecules or polypeptides, or their effectors or modulators, for manufacturing an agent for the diagnosis, prevention or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DP119; DP444; DP810; DP685; WE474; DP160; RA977; RA770; antidiabetic;
anorectic; neuroprotective; gene therapy; pancreatic disorder; diabetes;
neurodegenerative disorder; hyperglycaemia; impaired glucose tolerance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pancreatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dohrmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAY-2002;
17-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAY-2003; 2003WO-EP005700.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gallus gallus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chicken WE474 protein SEQ ID NO:28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADE43918;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE43918 standard; protein; 122 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (DEVE-) DEVELOGEN ENTWICKLUNGSBIOLOGISCHE FORSCH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2004-035045/03.
DB; ADE43917.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LGDIGD 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 28; 125pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disorders.
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2002EP-00020829.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Austen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            a Methanopyrus kandleri protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score
Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  non-human animal which over- or under-expresses the DP119, DP444, DP810 DP685, MEA774, DP160, RA977 or RA770 gene product. In addition, the nucleic acid molecule, polypeptide or an effector/modulator of the nucleic acid or polypeptide, is used in monitoring or controlling the function of a gene or gene product influenced or modified by a DP119, DP444, DP810, DP685, WE474, DP160, RA977 or RA770 polypeptide, or in identifying substances that interact with the polypeptide. The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay;
                                                                                                                                                                                                                                                               Skeiky YAW, Persing
L'maisonneuve J, Zha
                                                                                                                                                                                                                                                                                                                             21-APR-2000; 2000US-0199047P.
02-JUN-2000; 2000US-0208841P.
07-JUL-2000; 2000US-0216747P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Propionibacterium acnes immunogenic protein #24266.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAU63370 standard; protein; 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          beta-cells in vitro and/or in vivo.
                                                                                                                                                                                   Propionibacterium acnes vaccinating against and
                                                                                                                                                                                                                           N-PSDB; AAS59633.
                                                                                                                                                                                                                                                                                                                                                                                20-APR-2001; 2001WO-US012865.
                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200181581-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Propionibacterium acnes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-FEB-2002 (first entry)
                                                                                                                                                 Example 1;
                                                                                                                                                                                                                                                                                                     (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                       2001-616774/71.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           79
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                                                                                                                                                 SEQ ID NO 24565; 1069pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ₽
                                                                                                                                                                         vulgaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  osteopathic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           84
                                                                                                                                                                                                                                                                ing DH, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93.8%;
                                                                                                                                                                                   polypeptides and nucleic acids useful for diagnosing infections, especially useful ;
                                                                                                                                                                                                                                                                Mitcham (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 30; DB Pred. No. 3.8e 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              n vivo. (I) may also be used in preparing a or under-expresses the DP119, DP444, DP810,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     neuroprotectant.
                                                                                                                                                                                                                                                                JL, Wang S
, Carter D;
                                                                                                                                                  English
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0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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polypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO syndrome (synovitis, acne, pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central nervous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determined the contact that being a sample with a binding agent that binds to the proteins of the invention

Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic

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sample with a bi

amount of bound protein

in the sample.

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Query Match
Best Local S
Matches 5
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Zhang Y,
Barth B,
encoding a Propionibacterium acnes protein. The invention also relates to polypeptides encoded by the polynucleotides (ABM35624-ABM64536) and to immunogenic fragments of P. acnes polypeptides. The invention additionally encompasses expression vectors and host cells comprising a polynucleotide of the invention; antibodies against polypeptides of the invention; fusion proteins comprising a polypeptide of the invention; a method for stimulating an immune response specific for a P. acnes polypeptide and an isolated T cell population comprising T cells prepared via this method; a vaccine composition (comprising P. acnes polypeptides, antibodies, fusion proteins, T cell populations, or antigen-presenting cells that express the polypeptide); a method and kit for detecting or determining the presence or absence of P. acnes in a patient; and a method for inhibiting the development of P. acnes in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to downregulate expression and activity of P. acnes polypeptides and therefore treat P. acnes infections. The antibodies may also be used as diagnostic agents for determining P. acnes presence, for example, by enzyme linked immunosorbent assay (BLISA). Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                           New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                                                                                                                                     The invention relates to an isolated polynucleotide (ACF64435-ACF64733)
                                                                                                                                                                                                                                                                                          Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-381789/36.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Acne vulgaris; antiseborrhoeic; dermatological; antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Propionibacterium acnes predicted ORF-encoded polypeptide #24565.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABM59889 standard; protein; 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORIXA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wang S, Jen S, I
Vallieve-Douglass
                                                                                                                                                                                                                                                                                      SEQ ID NO 24565; 1481pp; English.
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ing S, Jen S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93.8%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Persing DH,
Lodes MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 30; DB Pred. No. 3.9e
1; Mismatches
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. 3.9e+02;
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Benson DR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Maisonneuve JL;
Jones R, Carter
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RESULT 39
ABM65515
ABM65515
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ABM65515
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mitcham JL,
Zhang Y, Wa
Barth B, Va
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  patient. The P. acnes polypeptides, polynucleotides, antibodies, fusion proteins. T cell populations or antigen-presenting cells that express the polypeptides are useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein. The polynucleotides can also be used as probes or primers for nucleic acid hybridiaation. The vaccine composition is useful for the stimulation of an immune response against P. acnes, or for treating acne, and the kit is useful for performing a diagnostic assay. The present sequence represents a polypeptide predicted to be encoded by an ORF (open reading frame) contained within the P. acnes polynucleotides of the
                         The invention relates to an isolated polynucleotide (ACF64435-ACF64733) encoding a Propionibacterium acnes protein. The invention also relates to polypeptides encoded by the polynucleotides (ABM35624-ABM64536) and to immunogenic fragments of P. acnes polypeptides. The invention additionally encompasses expression vectors and host cells comprising a polynucleotide of the invention; antibodies against polypeptides of the invention; a method for stimulating an immune response specific for a P. acnes polypeptide and an isolated T cell population comprising T cells prepared via this method; a vaccine composition (comprising P. acnes polypeptides,
                                                                                                                                                                                                                                                                                                                                                                                                                                New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
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                                                                                                                                                                                                                                                                                                                                                                              Claim 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-381789/36.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Acne vulgaris; antiseborrhoeic; dermatological; antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABM65515
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunostimulant;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           86
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                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 30191; 1481pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IGDLGD 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wang S, Jen S, I
Vallieve-Douglass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Skeiky YAW,
ang S, Jen S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immune response; vaccine; immunogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93.8%;
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Lodes MJ,
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Pred. No.
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Benson DR,
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Jones R, Carter
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polynucleotides,

antibodies,

fusion

proteins,

populations,

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RESULT 40
AAU65828
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Best Local
polypeptides. The the treatment, I P. acnes. The d
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay; dermatological; osteopathic; neuroprotectant.
                                                                                                                                                                                                                                                                                                                                      WPI; 2001-616774/71
                                                                                                                                                                                                                                                                                                                                                                                       Skeiky YAW, Persing DH, L'maisonneuve J, Zhang
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02-JUN-2000; 2000US-0208841P.
07-JUL-2000; 2000US-0216747P.
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                                                                                                                                                                                                                                                  Propionibacterium acnes
                                                                                                                                    Example 1; SEQ ID NO 27023; 1069pp; English.
                                                                                                                                                                                                                           nating
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                         AAU39105-AAU68017 represent Propionibacterium acnes immunogenic des. The proteins and their associated DNA sequences are used in ment, prevention and diagnosis of medical conditions caused by
                                                                                                                                                                                                                        against
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                                                                                                                                                                                                                        polypeptides and nucleic acids useful for diagnosing infections, especially useful
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                                                                                                                                                                                                                                                                                                                                                                                 Mitcham JL, Wang S;
, Jen S, Carter D;
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3.9e+02;
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밁 Ś

syndrome

encoding a Propionibacterium acnes protein. The invention also relates to polypeptides encoded by the polymucleotides (ABM35624-ABM64536) and to immunogenic fragments of P. acros polypeptides. The invention additionally encompasses expression vectors and host cells comprising a polymucleotide of the invention; antibodies against polypeptides of the invention; antibodies against polypeptides of the invention; a

The invention relates to an isolated polymucleotide (ACF64435-ACF64733) encoding a Propionibacterium acnes protein. The invention also relates

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Example 1; SEQ ID

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27023; 1481pp;

English.

protein.

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ABM62347
ID ABM62347
ID ABM62347
AC ABM62
AC ABM62
XX ACNE
DT 20-00
XX ACNE
KW ACNE
KW ACNE
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Cc P. acnes is also involved in infections of bone, joints and the central conervous system, however it is particularly involved in the inflammatory cc lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a cc sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The cc polypeptides may be used as antigens in the production of antibodies conspected for P. acnes proteins. These antibodies can be used to compregulate expression and activity of P. acnes polypeptides and cc devargelate expression and activity of P. acnes polypeptides and cc diagnostic agents for determining P. acnes presence, for example, by cc enzyme linked immunosorbent assay (ELISA). Note: The sequence data for this patent did not form part of the printed specification, but was cc for with a patent did not form the consence of the printed specification of the patent directly from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mitcham ,
Zhang Y,
Barth B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                           New Propionibacterium acnes polypeptides and polynucleotides encoding polypeptide, useful for diagnosing, preventing or treating acne vulgar or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-OCT-2002; 2002WO-US032727.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Propionibacterium acnes predicted ORF-encoded polypeptide #27023
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Propionibacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-OCT-2001; 2001US-00978825.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Acne vulgaris; antiseborrhoeic; dermatological; antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunostimulant;
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                                                                                                                                                                                                                                                                                                                                                                                             2003-381789/36.
DB; ACF64624.
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4; Conserv
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VGDVGD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard; protein; 261 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wang S, Jen S, Lod
Vallieve-Douglass J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Skeiky YAW,
ng S, Jen S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A
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Lodes MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 30; DB Pred. No. 8.2e 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bhatia
Benson D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ).2e+02;
0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 261;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Maisonneuve JL;
Jones R, Carte
                                                                                                                                                                                                                                                                                                              acne vulgaris,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Carter D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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RESULT 42
ABG14312
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Best Local
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                                                                                   New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutation responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                   WPI; 2001-639362/73.
N-PSDB; AAS78499.
   Claim 20;
                                                                                                                                                                                                                                                                                                                                                                                                                             31-MAR-2000; 2000US-00540217.
23-AUG-2000; 2000US-00649167.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-MAR-2001; 2001WO-US008631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel human diagnostic protein #14303.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABG14312 standard; protein; 275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 261 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200175067-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                       (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; chromosome mapping; gene mapping; gene therapy;
supplement; medical imaging; diagnostic; genetic d
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21 VGDVGD 26
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                                                                                                                                                                                                                                                                                                          孲,
   SEQ ID
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   ö
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44671;
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                                                                                                                                                                                                                                                                                                          Tang
                                                                                                                                                                                                                                                                                                          TY
103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 30; DB 6;
Pred. No. 8.2e+02;
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                                                                                         mutations
o assess
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The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping,

Claim 18; Page 161; 193pp; German

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RESULT 43
ABB77414
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cc and in recombinant production of (II). The polynucleotides are also used ci in diagnostica as expressed sequence tags for identifying expressed companies. (I) is useful in gene therapy techniques to restore normal cuseful for generating antibodies against it, detecting or quantitating a copypeptide in tissue, as molecular weight markers and as a food copypeptide in tissue, as molecular weight markers and as a food copypeptide and its binding partners are useful in medical imaging cof sites expressing (II). (I) and (II) are useful for treating disorders complyapetide and polynucleotide sequences have applications in companies, forensics, gene mapping, identification of mutations companies, forensics, gene mapping, identification of mutations companies for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and camino acid sequences. ABG00010-ABG30377 represent novel human diagnostic patent did not appear in the printed specification, but was obtained in constraint of the invention. Note: The sequence data for this cate of the invention. Note: The sequence data for this cate of the invention of the sequence data for this cate of the invention.
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                                           New nucleic acid from Ashbya gossypii, useful, when modulated, for producing fine chemicals, e.g. amino acids, encodes membrane-construction or -transport proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-AUG-2003
19-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 275 AA;
                                                                                                                N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Eremothecium gossypii.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fungal membrane transport protein SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABB77414
                                                                                                                                                               Althoefer H,
                                                                                                                                                                                                                                15-SEP-2000; 2000DE-01046074
                                                                                                                                                                                                                                                                 13-SEP-2001; 2001WO-EP010575
                                                                                                                                                                                                                                                                                                     21-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
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                                                                                                                                                                                                                                                                                                                                    WO200222824-A2
                                                                                                                                                                                                 (BADI ) BASF
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                                                                                                                2002-383130/41.
DB; ABL58819.
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5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first entry)
                                                                                                                                                                 Revuelta Doval JL,
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                                                                                                                                                                                                                                                                                                                                                                     /note= "Xaa = unknown,
                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                       .302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93.8%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 30; DB 4;
Pred. No. 8.6e+02;
1; Mismatches (
                                                                                                                                                               Santos
                                                                                                                                                                                                                                                                                                                                                                     encoded by degenerate codons"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NO 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 275;
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RESULT 44
ANBE66144
ID ABP66
XX ABP66
XX ABP66
XX I9-NC
XX Bifid
XX Bifid
XX Bifid
XX Bifid
XX Ident
XX Ident
XX IO-UA
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an isolated nucleic acid (I) from Ashbya
                                                                                                                                                                                                                                                                       Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                       Novel polynucleotide comprising Bifidobacterium genome sequence useful as a probe or primer for detecting and/or identifying Bifidobacterium longum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-668397/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EP1227152-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bifidobacterium longum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bifidobacterium longum NCC2705; Bifidobacterium; bacterial; antidiarrheic; antibacterial; inhibitor of Salmonella; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bifidobacterium longum NCC2705 ORF amino acid sequence SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-NOV-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABP66144 standard; protein; 308 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NEST ) SOC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CO
                                                                                                                                                                                                                                                                                                                                               biological sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standardise OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LCDICD
                                                                                                                                                                                                                                                                   SEQ ID NO 888; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROD NESTLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Æ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   composition; pharmaceutical composition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                lactic acid bacterium; diarrhoea; pathogenic bacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93.8%;
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Pred. No. 9.5e
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 5;
9.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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The present invention describes a polynucleotide (I) comprising a sequence of a Bifidobacterium genome selected from the nucleotide sequences given in ABQ81842 and ABQ81843, or a sequence sixen in ABQ81842 and ABQ81843. Also described is a polynucleotide (II) encodinable and ABQ81843. Also described is a polynucleotide (II) encodinable for the present of the prese

encoding

The invention relates to an isolated polynucleotide (ACF64435-ACF64733) encoding a Propionibacterium acnes protein. The invention also relates polypeptides encoded by the polynucleotides (ABM35624-ABM64536) and to

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Claim 7;

SEQ ID NO 30600; 1481pp; English.

New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.

밁 5

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CC heterologous polypepide. [7] has antidiarrheic and antibacterial CC activities, and can be used as an inhibitor of Salmonella. [1] (which is C a probe) is useful for the detection and/or identification of Salmonella. [1] (which is C Bifidobacterium longum in a biological sample. A carrier containing the CC lactic acid bacterium Bifidobacterium longum NCC2705 (CNCM I-2618) can be CC used for preventing and/or treating diarrhoea brought about by pathogenic CC bacteria and/or rotavirus. The carrier is a food composition selected CC products, ice-creams, fermented cereal based products, milk based from milk, yogurt, curd, cheese, fermented milks, milk based fermented CC products, ice-creams, fermented cereal based products, milk based CC products, infant formula, pet food or a pharmaceutical composition CC supplement, wet oral supplement, dry tube feeding or wet tube feeding. CC [1] is useful in DNA arrays or chips to carry out analysis of the CC expression of the Bifidobacterium gene. ABG81844 to ABG81850 represent CC Bifidobacterium related nucleotide sequences given in the Sequence CC specification. N.B. The sequence data for this patent is not represented CC in the printed specification but is based on sequence information CC supplied by the European Patent Office
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM65924
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                         Mitcham JL, Sactions, Jen S, Zhang Y, Wang S, Jen S, Zhang Y, Wallieve-Douglass J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fusion protein, comprising a sequence selected from 1097 sequences given in ABP65258 to ABP66354 ligated in frame to a polynucleotide encoding a heterologous polypeptide. (I) has antidiarrheic and antibacterial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 308 AA;
                                                                                                                                                       WPI; 2003-381789/36.
                                                                                                                                                                                                                                                                                   15-OCT-2001; 2001US-00978825
                                                                                                                                                                                                                                                                                                                  11-OCT-2002; 2002WO-US032727.
                                                                                                                                                                                                                                                                                                                                                   24-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                WO2003033515-A1
                                                                                                                                                                                                                                                                                                                                                                                                            Propionibacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Acne vulgaris; antiseborrhoeic; dermatological; antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Propionibacterium acnes immunogenic polypeptide #30600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABM65924;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM65924 standard; protein; 353
                                                                                                                                                                                                                                                                                                                                                                                                                                               immunostimulant; immune
                                                                                                                                                                                                                                                   (CORI-)
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                                                                                                                                                                                                                                                     CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                               response;
                                                                                                                                                                                      Persing DH,
Lodes MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 30; DB Pred. No. 9.7e 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                 vaccine;
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9.7e+02;
0;
                                                                                                                                                                                                       Bhatia A,
Benson DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ımmunogenic
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                                                                                                                                                                                                       Maisonneuve JL;
Jones R, Carte
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                                                                                                                                                                                                       Carter D;
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RESULT 46
ABO71148
ID ABO71148
ID ABO71148
ID ABO71
AC AB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 4
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                                                                                                                                                                                                                                   Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-FEB-1998;
27-JUL-1998;
                                                                                            Disclosure; SEQ ID NO 19894; 455pp; English
                                                                                                                                                                                   pathological conditions resulting from bacterial
                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; ABD04719.
                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-615309/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rubenfield MJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GENO-) GENOME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 IGDIGD 6
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98US-0094190P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nolling J, Deloughery C,
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                                                                                                                                                                                             infection.
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prophylaxis and treatment of pathological conditions resulting from a bacterial infection, for evaluating a compound, such as a polypeptide, for the ability to bind a P. aeruginosa nucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs, including anti-P. aeruginosa drugs, as templates for recombinant production of P. aeruginosa drugs, as templates for recombinant components for diagnosis and/or treatment of P. aeruginosa-caused infection, and in detection of P. aeruginosa sequences or other sequences of Pseudomonas species using biochip technology. Sequences ABO67826-ABO84396 represent P. aeruginosa polypeptides of the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format from USPTO at
Sequence 456 AA;
                                                                                                                                                                                                                                                                                                                                                                                                          polynucleotides encoding them. The therapy of pathological conditions,
                                                   seqdata.uspto.gov/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                The sequences are useful in ons, as molecular targets for
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Matches
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410 IGDLGD 415
        1 IGDIGD
                              Similarity
5; Conserv
                               Conservative
                                       93.8%;
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Pred. No. 1
                               Mismatches
                                        DB 7;
1.5e+03;
                                               Length 456
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RESULT 47
AD036971
ID AD0366
XX AD0366
XX DT 07-OC
XX Cell
XX Cell
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XX WO200
XX WO200
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XX WO201
XX WO201
XX WO21
XX WO21
XX WO21
XX WO21
XX WFI;
XX Clair
XX Clair Cell proliferation-related polypeptide #63. 07-OCT-2004 (first entry) ADQ36971; ADQ36971 standard; protein; 500

cell proliferation related polypeptide; differentiation; stress response cell proliferation; senescence;

Oryza sativa 22-JUL-2004. WO2004061122-A2.

26-DEC-2002; 2002US-0436565P 23-DEC-2003; 2003WO-US041200

(SYGN) SYNGENTA PARTICIPATIONS ત G

WPI; 2004-534388/51.

New nucleic acid molecule encoding a cell proliferation-related polypeptide, useful for modulating cell proliferation, senescence, differentiation, development, and stress response in plants, and for producing enhanced food crops.

Claim 28; SEQ ID NO 130; 408pp; English.

encoding a cell proliferation-related polypeptide. The nucleic acid molecule and the encoded polypeptide, and methods are useful for modulating cell proliferation, senescence, differentiation, development, and stress response in plants, and for producing enhanced food crops. The present sequence represents a cell proliferation-related polypeptide. The present sequence is published separately from the main body of the specification present invention relates to an isolated nucleic acid molecule EPO

Sequence 500 AA;

The invention relates to

Pseudomonas aeruginosa polypeptides and the

Length 500;

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                                                                                                                           The invention comprises the amino acid and coding sequences of rice stress-related proteins. The DNA and protein sequences of the invention are useful for regulating and controlling plant maturation and development, including proliferation, seenscence, disease-resistance, o stress response. They are also useful for producing transgenic plants having improved properties, e.g. tolerance to pests, herbicides, or biotic or abiotic stresses, improved nutritional value, increased yield or proliferation, or improved structure causing less loss from lodging shattering. The present amino acid sequence represents a rice stress-related protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rice; stress-related protein; plant maturation; plant development; plant proliferation; plant senescence; plant disease-resistance; plant stress response; transgenic plant; pest tolerance; herbicide tolerance; biotic stress tolerance; abiotic stress tolerance; improved nutritional value; increased yield; increased proliferation.
                                                                                                     Sequence 500 AA;
                                                                                                                                                                                                                                                                                    Claim 28;
                                                                                                                                                                                                                                                                                                                        New isolated nucleic acids and proteins, useful for producing transgenic plants having improved properties, e.g. tolerance to pests, herbicides, or biotic or abiotic stresses, improved nutritional value, or increased
                                                                                                                                                                                                                                                                                                                                                                                                                         Cooper B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-DEC-2003; 2003WO-US041098.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-JUL-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2004061080-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rice stress-related protein #25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-OCT-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADQ15641;
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)B; ADQ15640.
115
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                      1 IGDIGD 6
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VGDVGD 120
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                                                                                                                                                                                                                                                                                    ID NO 50;
                                                               93.8%;
                                                                                                                                                                                                                                                                                    551pp; English.
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Pred. No. 1.6e+03;
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Pred.
                                                    Mismatches
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                                                               30;
                                                               1.6e+03;
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RESULT

Query Match Best Local Similarity

93.8%;

Score Pred.

No;

1.6e+03; DB 8; B

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cc acid molecule containing a nucleic acid sequence containing the nucleic acid sequence containing an uncleic acid sequence conding the novel amylo cell containing the nucleic acid sequence; a cell containing the nucleic acid sequence; biological tissue containing the nucleic acid sequence; a foodstuff, a food additive or a modifier of foodstuff containing the nucleic acid sequence; a foodstuff, a food additive or a modifier of foodstuff containing the nucleic acid sequence; a computer readable recording ce encoding the mino acid sequence of the nucleic acid sequence ce encoding the amino acid sequence of the novel amylo maltase polypeptide; and a cyclic glucan obtained by reacting the novel amylo maltase polypeptide; condictiffs such as cooked rice, Japanese confectionery, snacks, bakeries, noodles, Chinese dumpling, shao-mai, fishery kneading goods, crefrigerating foods, baby foods, pet foods, pet foods, pet foods supplements, where the method involves adding the novel amylo maltase polypeptide to the foodstuff before or immediately after heat concessing of the raw material, where the novel amylo maltase polypeptide is useful for manufacturing glucan which has a cyclic glucan from the starch of the foodstuff. The novel amylo maltase polypeptide is useful for manufacturing glucan which has a cyclic structure of (alpha)-1,4-glycoside linkage, foodstuff materials and food additives, modifier of foodscuffs, a drink or eatable composition. Infinsion solution or composition for adhesion where the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          snack; noodle; Chinese dumpling; shao-mai; fishery kneading; refrigerated; baby food; pet; animal feed; drink; food supplement; cyclic glucan; Synechocystis sp. PCC 6803.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel amylo maltase polypeptide comprising amino acid sequence of a wild-type amylo maltase with a substitution, addition or deletion at a position, where the amino acid residue interacts with acarbose of amino acids other than the wild-type amylo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New mutated amylo maltase polypeptide having increased enzyme activity and reduced hydrolyzing activity, useful for manufacturing foodstuffs such as Japanese confectionery, noodles, baby foods, and food additives.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADQ15026 standard; protein; 505
                                                                  method involves reacting the saccharide containing the linear structure of (alpha)-1,4-glycoside linkage or its derivatives with the novel amyle maltase polypeptide. This sequence represents a Synechocystis sp. PCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-528719/51.
N-PSDB; ADQ15025.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-NOV-2002; 2002JP-00342966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synechocystis sp. PCC 6803
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Amylo maltase; transgenic; foodstuff; food additive; modifier;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synechocystis sp. PCC 6803 amylo maltase polypeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 maltase amino acid sequence. The invention further comprises: a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 9; SEQ ID NO 17; 51pp; Japanese
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(NIDE ) NEC CORP
                                                                                                                                         composition, infusion solution or composition for adhesion, where the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EZAKI GLICO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2003JP-00322319
                                               polypeptide
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Search completed: July 27, 2005, 01:17:29 Job time: 167 secs
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Matches 4
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                                                                                                                                                                                  The invention describes a new isolated nucleic acid encoding a Klebsiella pneumoniae polypeptide. Also described are: a recombinant expression vector comprising the nucleic acid, operably linked to a transcription regulatory element; and a cell comprising the recombinant expression vector. The nucleic acid is useful for preparing a vaccine composition against Klebsiella pneumoniae. This is the amino acid sequence of a Klebsiella pneumoniae polypeptide of the invention
                                                                                                                                                                                                                                                                                                                     New nucleic acid encoding a Klebsiella pneumoniae polypeptide, useful for preparing a vaccine composition against Klebsiella pneumoniae.
                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-895346/82.
N-PSDB; ACH98063.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Klebsiella pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABO64512 standard; protein; 544 AA.
                                                                                                                                                            Sequence 544 AA;
                                                                                                                                                                                                                                                                                                                                                                                                     Breton GL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US6610836-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Recombinant expression vector; transcription regulatory element; Klebsiella pneumoniae protein; antibacterial; Vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Klebsiella pneumoniae polypeptide seqid 11029.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AB064512;
                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 11029; 932pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-JAN-2000; 2000US-00489039.
                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENO-) GENOME THERAPEUTICS CORP.
                                                                                                    y Match 93.8%;
Local Similarity 66.7%;
hes 4; Conservative
                                                    121 VGDVGD 126
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                                                                                                      Score 30; DB 7; Length 544; Pred. No. 1.7e+03; 2; Mismatches 0; Indels
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are greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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    GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd.
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US-09-92-540-10629
US-09-925-991A-11029
US-09-252-991A-30097
US-09-252-991A-2675
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Sequence 7, Appli
Sequence 5488, Ap
Sequence 5489, Ap
Sequence 4, Appli
Sequence 1, Appli
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Sequence 16239, A
Sequence 10934, A
Sequence 11029, A
Sequence 17290, A
Sequence 30097, A
Sequence 30097, A
Sequence 30600, A
Sequence 2752, A
Sequence 2752, A
Sequence 26864, A
Sequence 27761, A
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US-09-583-110-3650	US-09-438-185A-764	US-09-134-001C-3918	US-09-614-912-82	US-09-198-452A-812	US-09-252-991A-23322	US-09-710-279-2786	US-09-710-279-2196	US-09-710-279-2044	US-09-710-279-1384	US-09-710-279-1220	US-09-710-279-1198	US-09-710-279-1152	US-09-710-279-458	US-09-710-279-156	US-09-710-279-148	US-09-614-912-74	US-09-614-912-62	US-09-902-540-10537	US-09-710-279-1234	US-09-710-279-2188	US-09-489-039A-7930	
Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	
3650, Ap	-	œ	≥		23322, A	2786, Ap	2196, Ap	2044, Ap	1384, Ap	1220, Ap	1198, Ap	1152, Ap	458, App	156, App	148, App	74, App1	•	10537, A	1234, Ap	2188, Ap	•	,

ALIGNMENTS

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; Sequence 20030, Application US/09248796A
; Patent NO. 6747137
; GENERAL INPORMATION:
APPLICANT: Keith Weinstock et al
; APPLICANT: Keith Weinstock et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES REL
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; CURRENT FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR PILING DATE: 1998-02-13
; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 20030
; LENGTH: 325
                                                                                                                                                                                                                                RESULT 2
US-08-470-179-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-248-796A-20030
                                                                                                  Sequence 7, Application US/08470179
Patent No. 5645994
GENERAL INFORMATION:
APPLICANT: Huang Ph.D, Wai Mun
TITLE OF INVENTION: Method and Compositions for
TITLE OF INVENTION: Identification of Species in a Sample
                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Candida albicans
                                                            NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Trask, Britt and STREET: P.O. Box 2550 CITY: Salt Lake City
                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                         299 ignigh 304
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Pred. No. 87;
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; Sequence 5489, Application US/09513999C
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Best Local Similarity
Watches 5; Conserve
               RESULT 4
US-09-513-999C-5489
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; ORGANISM: Homo sapiens
US-09-513-999C-5488
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                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
FILE REFERENCE: 59 US2.REG
                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 5488
                                                                                                                                                     Matches
                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 5488, Application US/09513999C Patent No. 6783961
                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 801-531-9168
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 59 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: FORIGINAL SOURCE:
ORGANISM: Ric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/470,179
FILING DATE:
CLASSIFICATION: 435
CCLASSIFICATION: 435
                                                                                                                                                                                                                                                                      ENGTH: 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Sweigert Ph.D, Susan E.
REGISTRATION NUMBER: 36,289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 801-532-1922
TELEFAX: 801-531-9168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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Pred. No. 22;
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Pred. No. 25;
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                                                                                                                                                                                 Length 69;
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GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclett, A.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
FITTE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
PALENT NO. 5783961
PILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOPTWARE: Patent.pm
SEQ ID NO 5489
RESULT 6
US-09-248-796A-21116
US-09-248-796A-21116
; Sequence 21116, Application US/09248796A
; Sequence 21116, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstock et al
; APPLICANT: Keith Weinstock et al
; TITLE OF INVENTION: NUCLEIC ACID AMD AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                       GRGANISM: Myxococcus xanthus
FEATURE:
KAME/KEY: unsure
LOCATION: (1)..(185)
GTHER INFORMATION: unsure at all Xaa locations
US-09-902-540-14654
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(1.5849)B
CURRENT APPLICATION NUMBER: U$01-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 14654
LENGTH: 185
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Best Local Similarity
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Matches 5; Conserv
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TYPE: PRT
                                                                                                                                                                                                                                                                62 VGDIGD
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                                                                                                                                                                                                                                                                                                                                                                             Score 31; DB Pred. No. 74;
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RESULT 7
US-08-631-607-4
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US-09-248-796A-21116
RESULT 8
US-09-098-358B-4
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                                                                                                                                                                                                    ; MOLECULE TYPE: protein US-08-631-607-4
                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/631,607
FILING DATE: 08-APR-1996
CLASSIFICATION: 514
ATTORNEY/AGPNT
                                                                                                                     Query Match
Best Local Similarity
Watches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4, Application US/08631607 Patent No. 5767252
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CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION: APPLICANT: Worley
                                                                                                                                                                                                                                                                                    TELEFAX: 619/678-5099
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 211 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 209
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Haile, Liea A.
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 072
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Worley et al, Paul
TITLE OF INVENTION: NOVEL NEU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
ADDRESSEE: Fish & R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: not
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 4225 EXCITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 96.9%;
Local Similarity 83.3%;
es 5; Conservative
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                                                                   151 VGDIGD 156
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                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                    not relevant
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                                                                                                                                                   96.9%;
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                                                                                                                                                   Score 31; DB 1;
Pred. No. 85;
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                                                                                                                                                                    Length 211;
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Sequence 1, Application US/10638333
Patent No. 6825020
GENERAL INFORMATION:
APPLICANT: Sakaguchi, No. 6825020uo
APPLICANT: Kuwahara, Kazuhiko
TITLE OF INVENTION: GANP Protein
FILE REFERENCE: 050208-0014
CURRENT APPLICATION NUMBER: US/10/638,333
CURRENT FILING DATE: 2003-08-12
PRIOR APPLICATION NUMBER: PCT/JP99/04634
                                                                                                                                                                                                                                                                                                         Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Sakaguchi, No. 6673913uo
APPLICANT: Kuwahara, Kazuhiko
TITLE OF INVENTION: GANP Protein
PILE REFERENCE: 050208-0014
CURRENT APPLICATION UMBER: U9/09/914,272A
CURRENT FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: PCT/JP99/04634
PRIOR FILING DATE: 1999-08-27
PRIOR PTLING DATE: 1999-08-27
PRIOR PTLING DATE: 1999-02-24
NUMBER: 05 SEQ ID NOS: 9
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US-09-098-358B-4
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LENGTH: 211
TYPE: PRT
ORGANISM: ARTIFICIAL
FEATURE:
                                                                                                                                                                                          RESULT 10
US-10-638-333-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn version 3.1 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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Best Local Similarity
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TYPE: PRT
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Local Similarity 83.3%;
nes 5; Conservation
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83.3%;
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Pred. No.
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Pred. No. 9.5e+02;
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LENGTH: 1971
TYPE: PRT
ORGANISM: Mouse
US-10-638-333-1
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US-09-902-540-16239
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15849)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 16239
LENGTH: 365
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Best Local Similarity
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                                                      NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 10934
LENGTH: 423
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ORGANISM: Myxococcus xanthus -09-902-540-10934
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
APPLICANT: Wiegand, Roger C.
APPLICANT: Wiegand, Myxococcus xanthus Genome Sequences and Uses Thereof FILE REPERENCE: 38-10(15849)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
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PRIOR APPLICATION NUMBER: 47035/1999
PRIOR FILING DATE: 1999-02-24
NUMBER OF SEQ ID NOS: 9
                                                                                                                PRIOR APPLICATION NUMBER: 60/217,883 PRIOR FILING DATE: 2000-07-10
                                       TYPE: PRT
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Pred. No. 2.4e+02;
2; Mismatches 0
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Pred. No. 9.5e+02;
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RESULT 15
US-09-252-991A-30097
; Sequence 30097, Application US/09252991A
; Patent No. 6551795
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Patent No. 6610836
GENERAL INFORMATION:
APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
FILE REFERENCE: 2709.2004001
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 11029
LENGTH: 544
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PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 19894
LENGTH: 456
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GENERAL INFORMATION:
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Best Local
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CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
                                                                                                                                                                                                                                                                            TYPE: PRT
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121 VGDVGD 126
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Pred. No.
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Pred. No. 3.7e+02;
_____matches 0;
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Pred. No. 3.1e+02;
1; Mismatches 0;
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. 2.8e+02;
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; Sequence 20035, Application US/09252991A
; Patent No. 6551795
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Best Local Similarity 83.3
Thes 5; Conservative
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NOS: 33142
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR PRIOR DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
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CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
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ORGANISM: Pseudomonas aeruginosa
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Similarity 66.7%;
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    Mismatches

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Pred. No. 5.9e+02;
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Pred. No.
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APPLICANT: MAIC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO P
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 30600
LENGTH: 514
TYPE: PRT
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US-09-252-991A-30600
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CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-02-19
PRIOR FILING DATE: 1998-07-27
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SEQ ID NO 20035
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Best Local Similarity
Matches 4; Conserv
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SEQ ID NO 26752
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Best Local Similarity
 Matches
                                     Query Match
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
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                 Local Similarity
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66.7%;
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90.6%; Score 29; DB 66.7%; Pred. No. 5.5% tive 2; Mismatches
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Pred. No.
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Pred. No. 4.7e+02;
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3.5e+02;
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                                         DB 4;
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RESULT 22
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; NAME/KEY: UNSURE
; LOCATION: (18),(19)
; OTHER INFORMATION: Identity of amino acid at the above locations are unknown.
US-09-252-991A-26864
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APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 26864, Application US/09252991A Patent No. 6551795
                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 15691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. 6833447
GENERAL INFORMATION:
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                                                                                                                                           Matches
                                                                                                                                                          Best Local Similarity
                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                          APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15949)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
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                                                                                                                                                                                                                                                  TYPE: PRT
                                                                                                                                                                                                                             ORGANISM: Myxococcus xanthus
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Pred. No. 7.4e+02;
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GENERRAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: ABRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                 Sequence 41400, Application US/09270767 Patent No. 6703491 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 7652
LENGTH: 72
TYPE: PRT
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Patent No. 6610836
GENERAL INFORMATION:
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LENGTH: 739
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPBUTICS
FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: US/09/489,039A
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and prof
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
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PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
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CURRENT FILING DATE: 1999-02-18
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Pred. No.
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Pred. No.
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                                                              of Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                                       Length 72
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NUMBER OF

ID NOS:

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FILLE OF ANTHOU-OTA
FILE REFERENCE: PATH00-07A
CURRENT APPLICATION NUMBER: US/09/583,110
CURRENT FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/107,433
PRIOR FILING DATE: 1998-06-30
PRIOR APPLICATION NUMBER: US 60/085,131
PRIOR FILING DATE: 1998-05-12
PRIOR PRILING DATE: 1998-05-12
PRIOR PILING DATE: 1997-07-02
PRIOR FILING DATE: 1997-07-02
NUMBER OF SEQ ID NOS: 5322
SEQ ID NO 4901
LENGTH: 137
TWYDE DRT
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US-09-270-767-56616
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US-09-583-110-4901
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SEQ ID NO 41400
LENGTH: 96
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 56616
LENGTH: 96
TYPE: PRT
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APPLICANT: Homburger et al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 56616, Application US/09270767
Patent No. 6703491
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                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4901, Application US/09583110 Patent No. 6699703
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
Query Match
Best Local Similarity
                                                       ORGANISM: Streptococcus pneumoniae -09-583-110-4901
                                                                                                                                                                                                                                                                                                                            APPLICANT: Lynn Doucette-Stamm et al.
TITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Relating to Streptococcus
TITLE OF INVENTION: Pneumoniae for Diagnostics and Therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Xaa means any amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Drosophila melanogaster
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  33 LGDLGD 38
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66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               87.5%;
66.7%;
 87.5%;
83.3%;
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Pred. No. 1.4e+02;
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Pred. No. 1.4e+02;
2; Mismatches 0
 Score
Pred.
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 No.
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                   Length 137;
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RESULT 28
US-09-328-352-7997
; Sequence 7997, Application US/09328352
; Patent No. 6562958
; GENERAL INFORMATION:
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Patent No. 6800744

Patent INFORMATION:
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
APPLICANT: Lynn A Doucette ACID AND AMINO ACID
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID
SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO: 5189:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/107,433
FILING DATE: 30-Jun-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/ 085131
FILING DATE: May 12, 1998
APPLICATION NUMBER: 60/051553
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-011
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
LOCATION: (B) LOCATION 1...142
SEQUENCE DESCRIPTION: SEQ ID NO: 5189:
                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein HYPOTHETICAL: YES ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: <Unknown>
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
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                                                                                                                                        57 IGDIGE
                                                                                                                                                                            1 IGDIGD 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 142 amino acids TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (781)893-5007
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Pred. No. 2.2e+02;
1; Mismatches 0
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; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: amino acid sequence US-09-710-279-2188
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US-09-489-039A-7930
                                                                                                                                                                                                                                                                                                                                                                                         US-09-710-279-2188
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                                                                                                                                                                                                                                                                                                                              Sequence 2188, Application US/09710279
Patent No. 6703492
GENERAL INFORMATION:
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Patent No. 6610836

GENERAL INFORMATION:
APPLICANT: Gary Breton et. al
APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PREUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 2709.2004001
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Best Local Similarity
Matches 5; Conserv
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SEQ ID NO 7930
LENGTH: 151
TYPE: PRT
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SEQ ID NO 7997
LENGTH: 146
TYPE: PRT
                                                                                                                 SEQ ID NO 2188
LENGTH: 198
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Gary L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PUJ480US
CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/164,258 PRIOR FILING DATE: 1999-11-09
                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                           NUMBER OF SEQ ID NOS: 4472
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                                                                        ORGANISM: Artificial Sequence
                                                                                                TYPE: PRT
                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
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5; Conserv
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100.0%; Pr
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 4; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.3e+02;
hes 0;
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US-09-902-540-10537; Sequence 10537, Application US/09902540; Patent No. 6833447
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                                                                                                                     ; ORGANISM: Myxococcus xanthus US-09-902-540-10537
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                                                                                                                                                APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15849)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 1637
LENGTH: 226
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1234
LENGTH: 223
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GENERAL INFORMATION:
APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS
FILB REFERENCE: PU3480US
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Best Local
                                                          Matches
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ORGANISM: Artificial Sequence
FEATURE:
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                                                         Local Similarity
nes 4; Conserv
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nes 5; Conserv
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83.3%;
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66.7%;
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Pred. No.
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Pred. No.
                                                         Mismatches
                                                                         3.6e+02;
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US-09-614-912-62

Application US/09614912

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APPLICANT: Lee, Jan. ....

APPLICANT: Sakai, Hajime
APPLICANT: Sakai, Perry
APPLICANT: Weng, Zude
APPLICANT: Weng, Tide
APPLICANT: Anderson, Shawn
TITLE OF INVENTION: Plant Metabolism Genes
FILE REFERENCE: BB1378 US NA
CURRENT APPLICATION NUMBER: US/09/614,912
CURRENT FILING DATE: 1299-07-12
PRIOR APPLICATION NUMBER: 60/143,401
PRIOR APPLICATION NUMBER: 60/143,401
PRIOR APPLICATION NUMBER: 60/143,401
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No. 6677502
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                                                                                                                                                                                                                                                         Patent No.
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TITLE OF INVENTION: Plant Metabolism Genes
FILE REFERENCE: BB1378 US NA
CURRENT APPLICATION NUMBER: US/09/614,912
CURRENT FILING DATE: 2000-07-12
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TYPE: PRT
ORGANISM: Zea mays
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NAME/KEY: UNSURE
LOCATION: (123)
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Ta, Zude
                                                                                                                                                                                                                                                                                                                                             GDIGD 78
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Rafalski,
                                                                                                                                                                                   Rafalski, Antoni
Orozco, Buddy
Miao, Gou-Hau
                                                                                                                                                                                                                                                                   Application US/09614912
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US-09-710-279-148
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TYPE: PRT
; ORGANISM: Triticum aestivum
US-09-614-912-74
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US-09-710-279-156
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                             Sequence 156, Application US/09710279

Patent NO. 6703492

GENERAL INFORMATION:
APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERWIDIS NUCLEIC ACIDS
FILE REFERENCE: PU3480US
CUCRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 1090-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver.
SEQ ID NO 148
LENGTH: 337
TYPE: PRT
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Patent No. 6703492
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Best Local (
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PRIOR FILING DATE: 1999-12-21
NUMBER OF SEQ ID NOS: 204
SOFTWARE: Microsoft Office of
                                                                                                                                                                                                                                                                                                                                                      Matches
SOFTWARE: Pa
SEQ ID NO 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
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PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence FEATURE:
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les 5; Conserv
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les 5; Conserv
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100.0%;
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83.3%;
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Pred. No. 5.
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5.5e+02;
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                                                                                                                                      AND PROTEINS
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CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 458
LENGTH: 337
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence; FEATURE:
THER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: amino acid sequence
US-09-710-279-458
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US-09-710-279-1152
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                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: KIMMERLY, WILLIAM JOHN
APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS
FILE REFERENCE: PU3480US
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 1152
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   Query Match
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Best Local Similarity
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TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: PU3480US
                                     FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthetic
OTHER INFORMATION: amino acid sequence
                                                                                                      ORGANISM: Artificial Sequence
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                                                                                                                                 TYPE: PRT
                                                                                                                                                ENGTH: 337
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83.3%;
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Pred. No. 5.5e+02;
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US-09-710-279-1198
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US-09-710-279-1220
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Matches
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1220
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1198
LENGTH: 337
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Patent No. 6703492
GENERAL INFORMATION:
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Matches 5; Conserv
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TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR PILING DATE: 1999-11-09
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CURRENT FILING DATE: 2000-11-09
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TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: amino acid sequence
                                                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
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Pred. No. 5.5e
1; Mismatches
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Pred. No. 5.5e
1; Mismatches
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5.5e+02
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RESULT

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RESULT 43
US-09-710-279-2196
; Sequence 2196, Application US/09710279
; Patent No. 6703492
; GENERAL INFORMATION:
; APPLICANT: KIMMERLY, WILLIAM JOHN
; TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS
; FILE REFERENCE: PU3480US
; CURRENT APPLICATION NUMBER: US/09/710,279
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Matches 5
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Best Local Similarity 83.2
Best Conservative
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SEQ ID NO 2044
LENGTH: 337
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Patent No. 6703492
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CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: Patentin Ver. 2.1
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TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR PILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
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TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: amino acid sequence
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nes 5; Conserv
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83.3%;
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Pred. No.
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Pred. No. 5.5e+02
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US-09-710-279-2786
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SEQ ID NO 23322
LENGTH: 353
TYPE: PRT
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TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 2786
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LENGTH: 337
                                                                                                                                                                                                                                                          Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J.
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Matches
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Best Local Similarity
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                                                                                                            APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 199-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,788
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SOFTWARE: PatentIn Ver. 2.1
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                                                                      PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER CENTROL DATE: 1998-07-27
                                                              NUMBER OF SEQ ID NOS:
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83.3%;
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83.3%;
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Pred. No. 5.5e+02;
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Pred. No. 5.
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5.5e+02;
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RESULT 46
US-09-198-452A-812
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                                           APPLICANT: Anderson, Shawn
TITLE OF INVENTION: Plant Metabolism Genes
FILE REFERENCE: BB1378 US NA
CURRENT APPLICATION NUMBER: US/09/614,912
CURRENT FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: 60/143,401
PRIOR APPLICATION NUMBER: 60/143,412
PRIOR APPLICATION NUMBER: 60/146,650
PRIOR FILING DATE: 1999-07-12
PRIOR APPLICATION NUMBER: 60/170,906
PRIOR FILING DATE: 1999-12-15
PRIOR APPLICATION NUMBER: 60/172,959
PRIOR APPLICATION NUMBER: 60/172,959
PRIOR FILING DATE: 1999-12-21
PRIOR APPLICATION NUMBER: 60/172,959
PRIOR APPLICATION NUMBER: 60/172,959
PRIOR APPLICATION NUMBER: 60/172,959
PRIOR APPLICATION NUMBER: 60/172,959
PRIOR APPLICATION NUMBER: 60/172,959
PRIOR APPLICATION NUMBER: 60/172,946
PRIOR APPLICATION NUMBER: 60/172,946
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US-09-614-912-82
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Patent No. 6599294
GENERAL INFORMATION:
APPLICANT: Griffais, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preventing of INVENTION; and treatment of infection
FILE REFERENCE: 9710-003-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 812
LENGTH: 361
TYPE: PRT
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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                   NUMBER OF SEQ ID NOS:
                                     PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
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Microsoft Office 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Allen, Ste
Rafalski,
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Lee, Jian Ming
Sakai, Hajime
                                                                                                                                                                                                                                                                                                                                                        Caimi, Perry G
                                                                                                                                                                                                                                                                                                                                                                              Sakai, Haj
Weng, Zude
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Orozco, Buddy
Miao, Gou-Hau
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/09614912
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                                   1999-12-21
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100.0%; Pred. No. -
100.0%; Mismatches
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100.0%; Pr
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; Pred. No.
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PRIOR APPLICATION NUMBER: US 60/108,279
PRIOR FILING DATE: 1998-11-12
PRIOR APPLICATION NUMBER: US 60/128,606
PRIOR FILING DATE: 1999-04-08
NUMBER OF SEQ ID NOS: 1074
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 764
LENGTH: 380
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Best Local Similarity
"---hes 5; Conserv:
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; TYPE: PRT
; ORGANISM: Oryza sativa
US-09-614-912-82
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US-09-134-001C-3918
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; OTHER INFORMATION: CPn0762
US-09-438-185A-764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:

APPLICANT: Lynn Doucette-Stamm et al
APPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: GTC-007
CURRENT APPLICATION NUMBER: US/09/134,001C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR APPLICATION NUMBER: US 60/064,764
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-11-08
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674

SEQ ID NO 3918
LENGTH: 375
                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Stephens, Richard
APPLICANT: Mitchell, Wayne
APPLICANT: Kalman, Sue
APPLICANT: Davis, Ronald
APPLICANT: Davis, Ronald
APPLICANT: The Regents of the University of California
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Chlamydia Pneumoniae Genome Sequence
FILE REFERENCE: 018941-000411US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3918, Application US/09134001C Patent No. 6380370
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LENGTH: 372
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No. 682207
                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/438,185A CURRENT FILING DATE: 2002-03-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Staphylococcus epidermidis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
                                      ORGANISM: Chlamydia pneumoniae FEATURE:
                                                                                TYPE: PRT
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83.3%;
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                                                                                                                                                                                                                                                                                                                               Pneumoniae Genome Sequence

    Mismatches

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Pred. No.
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GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al.
FITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Relating to Streptococcus
FITLE OF INVENTION: Pneumoniae for Diagnostics and Therapeutics
FILE REFERENCE: PATH00-07A
CURRENT FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/107,433
PRIOR FILING DATE: 1998-06-30
PRIOR APPLICATION NUMBER: US 60/085,131
PRIOR FILING DATE: 1998-05-12
PRIOR APPLICATION NUMBER: US 60/085,131
PRIOR FILING DATE: 1998-05-12
PRIOR APPLICATION NUMBER: US 60/085,131
PRIOR PILING DATE: 1997-07-02
NUMBER OF SEQ ID NOS: 5322
SEQ ID NO 3650
LENGTH: 388
TYPE: PRT
ORGANISM: Streptococcus pneumoniae
US-09-583-110-3650
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Search completed: July 27, 2005, 01:21:40 Job time : 24 secs
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US-09-583-110-3650
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                                                                                                                                                                                     Query Match 87.5%; Score 28; DB 4; Length 388; Best Local Similarity 100.0%; Pred. No. 6.4e+02; Matches 5; Conservative 0; Mismatches 0; Indels
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Database
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Maximum Match 100%
Listing first 50 summaries
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Maximum DB seq length: 200000000
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Perfect score:
                                                                                                                                                                                                           Published Applications AA:*

1: /cgn2 6/ptodata/2/pubpaa/US07 PUBCOMB.pep:*

2: /cgn2 6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

3: /cgn2 6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

4: /cgn2 6/ptodata/2/pubpaa/US06 PUBCOMB.pep:*

5: /cgn2 6/ptodata/2/pubpaa/US07 PUBCOMB.pep:*

6: /cgn2 6/ptodata/2/pubpaa/US08 NEW PUB.pep:*

7: /cgn2 6/ptodata/2/pubpaa/US08 NEW PUB.pep:*

9: /cgn2 6/ptodata/2/pubpaa/US08 PUBCOMB.pep:*

10: /cgn2 6/ptodata/2/pubpaa/US08 PUBCOMB.pep:*

11: /cgn2 6/ptodata/2/pubpaa/US08 PUBCOMB.pep:*

12: /cgn2 6/ptodata/2/pubpaa/US08 PUBCOMB.pep:*

13: /cgn2 6/ptodata/2/pubpaa/US08 PUBCOMB.pep:*

14: /cgn2 6/ptodata/2/pubpaa/US08 PUBCOMB.pep:*

15: /cgn2 6/ptodata/2/pubpaa/US10B.PUBCOMB.pep:*

16: /cgn2 6/ptodata/2/pubpaa/US10B.PUBCOMB.pep:*

16: /cgn2 6/ptodata/2/pubpaa/US10B.PUBCOMB.pep:*

16: /cgn2 6/ptodata/2/pubpaa/US10D.PUBCOMB.pep:*

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16: /cgn2 6/ptodata/2/pubpaa/US10D.PUBCOMB.pep:*
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32
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15.255 Million cell updates/sec
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Copyright (c) 1993 - 2005 Compugen Ltd.
/ Cgn2_6/ptodata/2/pubpaa/US10E_PUBCOMB.pep: *
/cgn2_6/ptodata/2/pubpaa/US10.NEW_PUB.pep: *
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/cgn2_6/ptodata/2/pubpaa/US11.NEW_PUB.pep: *
/cgn2_6/ptodata/2/pubpaa/US11.NEW_PUB.pep: *
/cgn2_6/ptodata/2/pubpaa/US60.NEW_PUB.pep: *
/cgn2_6/ptodata/2/pubpaa/US60_PUBCOMB.pep: *
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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US-10-027-015A-3 US-10-027-015A-4 US-10-027-015A-4 US-10-767-701-55899 US-10-369-493-18184 US-10-425-115-192813 US-10-767-701-36746 US-10-029-386-29095 US-10-425-115-238465 US-10-158-034-94 US-10-158-034-68 US-10-205-342-21	ID
Sequence 3, Appli Sequence 4, Appli Sequence 53899, A Sequence 18184, A Sequence 192813, Sequence 36746, A Sequence 29095, A Sequence 230465, Sequence 238465, Sequence 94, Appl Sequence 68, Appl Sequence 21, Appl	Description

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RESULT 1 US-10-027-015A-3

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APPLICANT: AKELLA, RAMA
APPLICANT: RANIERI, JOHN P.
APPLICANT: RANIERI, JOHN P.
APPLICANT: RANIERI, JOHN P.
APPLICANT: RANIERI JOHN P.
APPLICANTION: ANGIOCENESIS
FILE REFERENCE: 2103.00500
CURRENT APPLICATION NUMBER: US/10/027,015A
CURRENT FILING DATE: 2001-12-21
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin version 3.2
SEQ ID NO 3
                                                                         Query Match
Best Local S
Matches 6
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                                                                                                                                                                                                                                 LENGTH: 6
TYPE: PRT
                                                                                                                                                                                          ORGANISM: Artificial FEATURE:
                                                                                          Local Similarity
                                    ignign 6
                                                                         100.0%; Score 32; DB 18; ilarity 100.0%; Pred. No. 1.6e+06; Conservative 0; Mismatches 0;
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                                                                                                                  Length 6;
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Sequence 53899, Application US/10767701

Publication No. US20040172684A1

GENERAL INFORMATION:
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Vongwei
APPLICANT: Cao, Vongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement
FILE REFERENCE: 38-21(5353)B
CURRENT APPLICATION NUMBER: US/10/767,701
CURRENT FILING DATE: 2004-01-29
NUMBER OF SEQ ID NOS: 63128
SEQ ID NO 53899
LENGTH: 153
TYPE: PRT
ORCANISM: Sorghum bicolor
FEATURE:
CHIEFE INVENTION: Close ID 128668
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Publication No. US20050143312A1
GENERAL INFORMATION:
APPLICANT: AKELLA, RAMA
APPLICANT: RANIERI, JOHN P.
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 4
Sequence 18184, Application US/10369493
Publication No. US20030233675A1
GENERAL INFORMATION:
APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR PROMOTING MYOCARDIAL AND PERIPHERAL TITLE OF INVENTION: ANGIOGENESIS
FILE REFERENCE: 2103.000500
CURRENT APPLICATION NUMBER: US/10/027,015A
CURRENT FILING DATE: 2001-12-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial FEATURE:
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TYPE: PRT
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Pred. No. 1.6e+06;
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Pred. No. 1.3e+02;
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US-10-425-115-192813; Sequence 192813, Application US/10425115; Publication No. US20040214272A1
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SEQ ID NO 192813
LENGTH: 689
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SEQ ID NO 18184
LENGTH: 515
                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                           Sequence 36746, Application US/10767701 Publication No. US20040172684A1
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Best Local
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement
FILE REFERENCE: 38-21 (53535)B
CURRENT APPLICATION NUMBER: US/10/767,701
CURRENT FILING DATE: 2004-01-29
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CURRENT FILING DATE: 2003-04-28
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21 (53222)B
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NAME/FCN: unsure
LOCATION: (1)..(689)
OTHER INFORMATION: unsure at all Xaa locations
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Pred. No.
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Pred. No. 6.3e+02;
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US-10-425-115-238465, Application US/10425115; Sequence 238465, Application US/10425115; Publication No. US20040214272A1; GENERAL INFORMATION:
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APPLICANT: La ROSA, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules an
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21 (53222) B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
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SEQ ID NO 36746
LENGTH: 80
TYPE: PRT
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Best Local :
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CURRENT FILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
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APPLICANT: Rank, David R.
APPLICANT: Hank, David K.
APPLICANT: HOWNI GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
FILE REFERENCE: ABOMICA-X-2
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OTHER INFORMATION:
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N: EXPRESSED IN FETAL LIVER, SIGNAL = 1.2

N: EXPRESSED IN LUNG, SIGNAL = 1.4

N: EXPRESSED IN HELA, SIGNAL = 1.4

N: EXPRESSED IN BAAIN, SIGNAL = 2.1

N: EXPRESSED IN HEART, SIGNAL = 2.1

N: EXPRESSED IN ADDIT LIVER, SIGNAL = 3.1

N: EXPRESSED IN BONE MARROW, SIGNAL = 3.1
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Pred. No. 1e+02;
1; Mismatches
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Pred. No. 1e+02;
1; Mismatches
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                                                      RESULT 10
US-10-158-034-68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 94, Application US/10158034
Publication No. US20030219758A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
Sequence 68, Application US/10158034
Publication No. US20030219758A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. SEQ ID NO 94
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Best Local Similarity
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                                                                                                                                                                                                   Query Match
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OTHER INFORMATION: Xaa e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/158,034
CURRENT FILING DATE: 2002-05-31
CURTOR Application removed - See File Wrapper
NUMBER OF SEQ ID NOS: 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Nucleic Acids, Proteins, FILE REFERENCE: PTZ10C1
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TYPE: PRT
                                                                                                                                                                                                                                              OTHER INFORMATION: Xaa equals any
                                                                                                                                                                                                                                                              NAME/KEY: misc feature LOCATION: (128)
                                                                                                                                                                                                                                                                                                     LOCATION: (127)
OTHER INFORMATION: Xaa equals
                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (123)
OTHER INFORMATION: Xaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                LOCATION: (126)
OTHER INFORMATION: Xaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
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Pred. No. 1.7e+02;
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TITLE OF INVENTION: Nucleic Acids, Proteins, FILE REFERENCE: PTZ10C1
CURRENT APPLICATION NUMBER: US/10/158,034

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GENERAL INFORMATION:
APPLICANT: Schlegel, Robert
APPLICANT: Monahan, John E.
APPLICANT: Endege, Wilson O.
APPLICANT: Gannavarapu, Manji
APPLICANT: Gorbatcheva, Belli
APPLICANT: Hoersch, Sebastia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: PRT
; ORGANISM: Mus musculus
; FEATURE:
; OTHER INFORMATION: Protein: Cofilin
US-10-205-342-21
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US-10-205-823-72
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US-10-158-034-68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21, Application US/10205342
Publication No. US20030108906A1
GENERAL INFORMATION:
APPLICANT: Warner-Lambert Company
APPLICANT: Lee, Kevin
APPLICANT: Dixon, Alistair
APPLICANT: Brookebank, Robert
APPLICANT: Pinnock, Robert
TITLE OF INVENTION: Identification and Use of Molecules Implicated in Pain
FILE REFERENCE: WL-A-018198
CURRENT FILING DATE: 2002-07-24
PRIOR FILING DATE: 2002-07-24
PRIOR APPLICATION UNUBER: GB 0118354.0
PRIOR FILING DATE: 2001-07-27
PRIOR FILING DATE: 2001-07-27
PRIOR FILING DATE: 2001-07-27
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SOFTWARE: PATENTIN VERSION 3.0
SEQ ID NO 21
LENGTH: 166
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SEQ ID NO 68
LENGTH: 156
TYPE: PRT
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Best Local
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APPLICANT:
APPLICANT:
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APPLICANT:
                    TITLE
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                                      TITLE OF
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les 5; Conserv
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T: Anderson, Dustin
INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND
INVENTION: METHODS FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND
INVENTION: THERAPY OF PROSTATE CANCER
                                                                                                                                                                                                                                                                                                                                                                                                          1 IGDIGD 6
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                                                                                                      Hoersch, Sebastian
Kamatkar, Shubhangi
Wonsey, Angela M.
                                                                                                                                                         Gannavarapu, Manjula
Gorbatcheva, Bella
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Endege, Wilson O.
                                                                                    Wonsey, Angel
Glatt, Karen
                                                                     Zhao, Xumei
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Pred. No. 2.1e+02;
1; Mismatches 0;
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APPLICANT: Gibson, Bradford W.
APPLICANT: Taylor, Steven W.
APPLICANT: Glenn, Gary M.
APPLICANT: Glenn, Gary M.
APPLICANT: Warnock, Dale E.
APPLICANT: Warnock, Dale E.
TITLE OF INVENTION: TARGETS FOR THERAPEUTIC INTERVENTION
TITLE OF INVENTION: IDENTIFIED IN THE MITOCHONDRIAL PROTEOME
FILE REFERENCE: 660088.465
FILE REFERENCE: 620088.465
CURRENT APPLICATION NUMBER: US/10/408,765A
CURRENT FILING DATE: 2003-04-04
NUMBER OF SEQ ID NOS: 3077
NUMBER OF SEQ ID NOS: 3077
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                       Query Match
Best Local Similarity
Tatches 5; Conserve
US-10-425-115-256293, Application US/10425115; Sequence 256293, Application US/10425115; Publication No. US20040214272A1; GENERAL INFORMATION:
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CURRENT FILING DATE: 2002-07-25
PRIOR APPLICATION NUMBER: 60/307,982
PRIOR FILING DATE: 2001-07-25
PRIOR PILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/314,356
PRIOR PILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/325,020
PRIOR FILING DATE: 2001-09-25
PRIOR FILING DATE: 2001-09-25
PRIOR APPLICATION NUMBER: 60/341,746
PRIOR PILING DATE: 2001-12-12
PRIOR APPLICATION NUMBER: 60/362,158
PRIOR FILING DATE: 2002-03-05
PRIOR FILING DATE: 2002-03-05
PRIOR FILING DATE: 2002-03-05
PRIOR FILING DATE: 2002-03-05
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                                                                                 RESULT 14
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 72
LENGTH: 166
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APPLICANT: Fahy, Eoin D.
APPLICANT: Zhang, Bing
                                                                                                                                                                                                                                                                                                                                                 LENGTH: 16
TYPE: PRT
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Pred. No.
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Pred. No. 2.2e+02;
                                                                                                                                                                                                                            Mismatches
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei

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Query Match
Best Local Similarity
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US-10-224-951-4
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                                                                                                            APPLICANT: THE JOHN HOPKINS SCHOOL OF MEDICINE
APPLICANT: WORLEY, PAUL
APPLICANT: TSUI, CYNTHIA
TITLE OF INVENTION: NOVEL NEURONAL CELL GROWTH FAC
FILE REFERENCE: JHU 1340-1 (07265/086002)
CURRENT APPLICATION UNMBER: US/10/224,951
CURRENT FILING DATE: 2002-08-20
PRIOR FILING DATE: 2002-08-20
PRIOR FILING DATE: 1998-06-16
PRIOR FILING DATE: 1998-06-16
PRIOR FILING DATE: 1998-06-16
PRIOR FILING DATE: 1998-06-16
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CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 256293
                                                                                                                                                                                                                                                                         Sequence 4, Application US/10224951
Publication No. US20030022313A1
GENERAL INFORMATION:
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SEQ ID NO 93
LENGTH: 205
TYPE: PRT
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Best Local (
                                                  SOFTWARE: PatentIn version 3.0
SEQ ID NO 4
LENGTH: 211
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/158,034
CURRENT FILING DATE: 2002-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Rosen et al. TITLE OF INVENTION: Nucleic Acids, Proteins, FILE REFERENCE: PTZ10C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 176
TYPE: PRT
ORGANISM: Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
ORGANISM: Artificial Sequence FEATURE:
OTHER INFORMATION: RAT CRP
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5; Conserv
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No. US20030219758A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96.9%;
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83.3%;
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Pred. No. 2.4e
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.8e+02;
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2.4e+02;
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US-10-739-930-7499
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US-10-739-930-7500
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LENGTH: 299
TYPE: PRT
RGANISM: Zea mays
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Best Local
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                                   Matches
                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Zea mays
FEATURE:
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     1 IGDIGD 6
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; OTHER INFORMATION: Clone ID: US-10-739-930-7500
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APPLICANT: KOVALIC, DAVID K.
APPLICANT: KOVALIC, DAVID K.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH
TITLE OF INVENTION: PLANTS AND USES THEREOF FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53377)B
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SEQ ID NO 7500
LENGTH: 276
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                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 7499, Application US/10739930 Publication No. US20040216190A1
                                                                                                                                                                                                                                                                                                                                      APPLICANT: KOVALIC, DAVIÓ K.
TITLE OF INVENTION: NUCLBIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH
TITLE OF INVENTION: PLANTS AND USES THEREOF FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21 (53377) B
                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/739,930 CURRENT FILING DATE: 2003-12-18
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                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 11088
                                                                                                                   NAME/KEY: unsure
LOCATION: (1)..(299)
OTHER INFORMATION: unsure at all Xaa locations
                                                                                      FEATURE:
OTHER INFORMATION: Clone ID:
                 Local Similarity
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5; Conserv
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Conservative
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83.3%;
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                 96.9%;
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                 Score 31; DB 16;
Pred. No. 4.1e+02
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Pred: No. 2.9e+02;
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                 4.1e+02;
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3.8e+02;
ches 0;
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RESULT 19 US-10-156-761-11418

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APPLICANT: HATTORI, MASAHIRA
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-262
FILE REFERENCE: 249-262
CURRENT APPLICATION NUMBER: US/10/156,761
CURRENT FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: JP 2001-204089
PRIOR FILING DATE: 2001-05-30
PRIOR FILING DATE: 2001-05-30
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LENGTH: 328
TYPE: PRT
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SOFTWARE: PatentIn version
SEQ ID NO 367
LENGTH: 342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Publication No. US20040025202A1 GENERAL INFORMATION:
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/389,566
CURRENT FILING DATE: 2003-03-31
PRIOR APPLICATION NUMBER: US 60/365,301
PRIOR FILING DATE: 2002-03-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: JP 2001-272697 PRIOR FILING DATE: 2001-08-02
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APPLICANT: IKEDA, HARUO
APPLICANT: ISHIKAWA, JUN
APPLICANT: HORIKAWA, HIRO
                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/391,786
PRIOR FILING DATE: 2002-06-25
PRIOR APPLICATION NUMBER: US 60/392,018
PRIOR FILING DATE: 2002-06-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Laurie, Cathy C
TITLE OF INVENTION: Nucleic Acid Molecules Associated with Oil in Plants
FILE REFERENCE: 38-77(52900)D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Monsanto Technology, LLC APPLICANT: Laurie, Cathy C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 15109
                                                                                                             NAME/KEY: misc_feature
LOCATION: (1)..(1)
OTHER INFORMATION: Xaa can be any naturally occurring amino acid
                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Zea mays
                                            OTHER INFORMATION: Xaa
                                                              NAME/KEY: misc_feature
LOCATION: (10)..(10)
                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
                                                                                                         FEATURE:
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HORIKAWA, HIROSHI
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                                          can be any naturally occurring amino acid
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Pred. No. 4.5e
1; Mismatches
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4.5e+02;
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Query Match
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Watches 5; Conserve
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; OTHER INFORMATION: Xaa can be any naturally occurring amino acid US-10-389-566-367
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APPLICANT: Cao Yongwei

TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

FILE REFERENCE: 38-21(5323)B

CURRENT APPLICATION NUMBER: US/10/424,599

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 285684

SEQ ID NO 218355
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US-10-424-599-218355
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SEQ ID NO 23444
LENGTH: 364
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Publication No. US20030233675A1
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10 (52052) B.
FILE REFERENCE: 38-10 (52052) B.
                                                                                                                                                                                                                                                                                    APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/360,039 PRIOR FILING DATE: 2002-02-21
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CURRENT FILING DATE: 2003-02-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Deinococcus radiodurans
NAME/KEY: unsure
LOCATION: (1)...(464)
OTHER INFORMATION: unsure at all Xaa locations
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OTHER INFORMATION: Xaa can be any naturally occurring amino acid
                                                                 ORGANISM: Glycine max FEATURE:
                                                                                                        TYPE: PRT
                                                                                                                          LENGTH:
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83.3%;
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Pred. No.
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Pred. No.
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1 IGDIGD 6

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; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_39201C.1.pep
US-10-424-599-218355
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US-10-425-115-254563
                                                                                  ; ORGANISM: Mesorhizobium loti
US-10-369-493-11876
                                                                                                                                            TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES FILE REFERENCE: 38-10(52052)B
CURRENT APPLICATION UNMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR PRILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ.ID NO 11876
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LENGTH: 537
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Best Local Similarity
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              Query Match
Best Local Similarity
Matches 5; Conserv
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Zea mays
FEATURE:
                                                                                                                   LENGTH: 693
TYPE: PRT
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les 5; Conserva
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83.3%;
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Pred. No. 6.5e+02;
1; Mismatches 0
            Score 31; DB 15
Pred. No. 1e+03;
1; Mismatches
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Pred. No. 7.6e+02;
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                                                  DB 15; Length 693;
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RESULT 27
US-10-732-923-4243
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APPLICANT: LANGU Yihua

TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

FILE REFERENCE: 38-21(53223)B

CURRENT PILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 285684

SEQ ID NO 235305

LENGTH: 115

TYPE: PRT

ORGANIER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-424-599-235305; Sequence 235305, Application US/10424599; Publication No. US20040031072A1
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Best Local Similarity
"---hes 5; Conserv
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US-10-747-133-1
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APPLICANT: Kuwahara, Kazuhiko
TITLE OF INVENTION: GANP Protein
TITLE OF INVENTION: GANP Protein
TITLE REFERENCE: 050208-0014
CURRENT APPLICATION NUMBER: US/10/747,133
CURRENT FILING DATE: 2001-12-05
PRIOR APPLICATION NUMBER: US/09/914,272
PRIOR FILING DATE: 2001-12-05
PRIOR APPLICATION NUMBER: PCT/JP99/04634
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: 47035/1999
PRIOR FILING DATE: 1999-02-24
NUMBER OF SEQ ID NOS: 9
                                                                                                                                              US-10-424-599-235305
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                                                                                         Query Match
Best Local Similarity
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TYPE: PRT
ORGANISM: Mouse
                                                                                                                                                                                   ORGANISM: Glycine max FEATURE:
                                                                                                                                                                 OTHER INFORMATION: Clone ID: PAT_MRT3847_54508C.1.pep
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38 LGDIGD 43
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                                                                          Conservative
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83.3%;
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Pred. No. 3e+03;
1; Mismatches 0; Indels
                                                                                         Score 30; DB 15; Length 115; Pred. No. 2.4e+02;
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US-10-425-115-335683; Sequence 335683; Application US/10425115; Publication No. US20040214272A1; GENERAL INFORMATION:
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules an
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21 (53222) B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
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US-10-425-115-185817
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US-10-732-923-4243
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Matches
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APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 185817
LENGTH: 204
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APPLICANT: Edgerton, Michael D
TITLE OF INVENTION: TRANSGERIC PLANTS WITH IMPROVED PHENOTYPES
FILE REFERENCE: 38-15 (52796) C
CURRENT APPLICATION NUMBER: US/10/732,923
CURRENT FILING DATE: 2003-12-10
PRIOR APPLICATION NUMBER: 10/310,154
PRIOR APPLICATION NUMBER: 10/310,154
PRIOR FILING DATE: 2002-12-04
NUMBER OF SEQ ID NOS: 24149
SEQ ID NO 4243
LENGTH: 115
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ORGANISM: Zea mays
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les 5; Conserv
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Pred. No. 4.3e+02;
1; Mismatches 0
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Pred. No.
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                                                                                                        and Other Molecules Associated With
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US-10-425-114-68337; Sequence 68337, Application US/10425114; Publication No. US20040034888A1
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                              APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REPERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 68337
LENGTH: 477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local :
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LENGTH: 328
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Publication No. US20040034888A1
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LENGTH: 247
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APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5331)B
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
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APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, Dav
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APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, Dav
                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Zea mays
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ORGANISM: Zea mays
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nes 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                              243 VGDVGD
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Kovalic, David K.
Screen, Steven E
Tabaska, Jack E
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    Mismatches

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Pred. No.
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Pred. No. 7.2e+02
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules an
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 325328
LENGTH: 486
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US-10-425-114-57203
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LENGTH: 486
TYPE: PRT
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Best Local Similarity
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Best Local (
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CURRENT APPLICATION NUMBER: US/10/425,114

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 73128
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Zea mays FEATURE:
                      NAME/KEY: unsure
LOCATION: (1)..(486)
OTHER INFORMATION: unsure at all Xaa locations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Clone ID: UC-ZMFLB73152G04_FLI.pep
OTHER INFORMATION: Clone ID: MRT4577_59770C.1.pep
                                                                                                        ORGANISM: Zea mays
                    FEATURE:
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Cao, Yongwei
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83.3%;
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Pred. No. 1.1e+03;
2; Mismatches 0
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Pred. No.
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APPLICANT: CAOVALIC, David K.

APPLICANT: Chou, Yihua

APPLICANT: Chou, Yihua

ITITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

ITITLE OF INVENTION: Plants

ITITLE REPERENCE: 38-21 (5322)B

CURRENT APPLICATION NUMBER: US/10/425,115

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 369326

SEQ ID NO 239897

LENGTH: 540

TYPE: NO.
                                                              ; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_125408C.1.pep
US-10-424-599-170978
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Best Local S
Matches 5
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CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 170978
LENGTH: 567
Matches
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                             Query Match
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                                                                                                                                                                                                                                                                                                APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
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                                                                                                       NAME/KEY: unsure
LOCATION: (1)...(567)
OTHER INFORMATION: unsure at all Xaa locations
                                                                                                                                                                  FEATURE:
                                                                                                                                                                              TYPE: PRT
ORGANISM: Glycine max
             Local Similarity
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tes 4; Conserv
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               93.8%;
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               Score 30; DB 15;
Pred. No. 1.3e+03;
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                               DB 15; Length 567;
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1 IGDIGD 6

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; Sequence 139894, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
   APPLICANT: La Rosa, Thomas J.
   APPLICANT: Kovalic, David K.
   APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
   APPLICANT: Wu, Wei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         맑
                                                       ; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_41143C.1.pep
US-10-437-963-139894
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US-10-437-963-184415
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                                                                                                                                                                         TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)B CURRENT APPLICATION NUMBER: US/10/437,963 CURRENT FILING DATE: 2003-05-14 NUMBER OF SEQ ID NOS: 204966 SEQ ID NO 139894
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 184415
LENGTH: 568
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APPLICANT:
APPLICANT:
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CURRENT FILING DATE: 2003-05-14
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
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                                                                                                               TYPE: PRT
ORGANISM: Oryza sativa
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LOCATION: (1)..(568)
OTHER INFORMATION: u
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ORGANISM: Oryza sativa
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                                                                                                                                                      ENGTH: 598
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Barbazuk, Brad
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93.8%;
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Score
Pred.
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30;
DB 16;
1.3e+03;
                  Length 598;
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6, Application US/10937730A
Publication No. US20050112726A1
GENERAL INFORMATION:
APPLICANT: HU, Znihao
APPLICANT: REID, Ralph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)B CURRENT APPLICATION NUMBER: US/10/437,963 CURRENT FILING DATE: 2003-05-14 NUMBER OF SEQ ID NOS: 204966 SEQ ID NOS: 204966
                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 181573, Application US/10437963 Publication No. US20040123343A1
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Best Local Similarity
                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Biosynthetic Gene Cluster for Leptomycin FILE REFERENCE: 01092.02 CURRENT FILING DATE: 2004-09-08 PRIOR APPLICATION NUMBER: US/10/937,730A CURRENT FILING DATE: 2004-09-08 PRIOR APPLICATION NUMBER: 60/502,423 PRIOR FILING DATE: 2003-09-11 PRIOR PPLICATION NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER: 60/503,384 PRIOR FILING DATE: 2004-03-15 NUMBER:
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APPLICANT:
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ORGANISM: Oryza sativa
FEATURE:
OTHER INFORMATION: Clone ID: PAT_MRT4530_78833C.1.pep
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                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
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1930 VGDVGD 1935
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                                                                                                                                                               93.8%;
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66.7%;
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                                                                                                                                2; Mismatches
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Pred. No. 1.5e+03;
2; Mismatches (
                                                                                                                                                            Score 30; DB 17;
Pred. No. 8.4e+03;
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US-10-425-115-253571
Sequence 253571, Application US/10425115
Publication No. US20040214272A1
GENERAL INFORMATION:
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                                                                                                   US-10-424-599-263936
                                                                                                                   RESULT 42
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                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants
FILE REPERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 253571
LENGTH: 79
TYPE: PRT
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LENGTH: 73
TYPE: PRT
ORGANISM: Glycine max
Sequence 263936, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: La ROSa Thomas J
APPLICANT: Kovalic David K
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Best Local S
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Best Local
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5523)B
FULE REPERENCE: 38-21(5523)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 203-04-28
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
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                                                                                                                                                                                                                                                                    Score 29;
Pred. No.
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Pred. No. 2.3e+02;
2; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                       DB 16; Length 79;
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yinua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and
TITLE OF INVENTION: Plants
FILLE REPERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 355214
LENGTH: 152
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; OTHER INFORMATION: Clone ID: PAT_MRT3847_80353C.1.pep
US-10-424-599-263936
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; OTHER INFORMATION: Clone ID:
US-10-425-115-355214
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US-10-767-701-60220
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Best Local S
Matches 4
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SEQ ID NO 263936
LENGTH: 90
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                                                                                                                                 Sequence 60220, Application US/10767701
Publication No. US20040172684A1
GENERAL INFORMATION:
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Best Local
APPLICANT: Kovalic, David K.

APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement
FILE REFERENCE: 38-21(53535)B
CURRENT APPLICATION NUMBER: US/10/767,701
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CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
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APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
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ORGANISM: Glycine max
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LOCATION: (1)..(152)
OTHER INFORMATION: u
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ORGANISM: Zea mays
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5; Conserv
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Pred. No. 2.9e+02;
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Pred. No. 5e+02;
2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                     DB 16; Length 152;
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US-10-767-701-60220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT FILING DATE: 2004-01-29
NUMBER OF SEQ ID NOS: 63128
SEQ ID NO 60220
LENGTH: 174
TYPE: PRT
ORGANISM: Sorghum bicolor
                                                                                                                                                                                                                                        Sequence 201329, Application US/10425115
Publication No. US20040214272A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 146654
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 146654, Application US/10437963 Publication No. US20040123343A1
              NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 201329
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                                                                                              APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
                                                         CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                    APPLICANT: La Rosa, Thomas J. APPLICANT: Kovalic, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Oryza sativa
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ENGTH: 189
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Barbazuk, Brad
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Pred. No. 6.2e+02;
""" ematches 0;
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Pred. No.
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5.8e+02;
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TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 156021
LENGTH: 321
TYPE: PPT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_55729C.1.pep
US-10-437-963-156021
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US-10-437-963-102521
                  APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT PILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 102521
LENGTH: 355
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Matches 5; Conserv
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Publication No. US20040123343A1
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APPLICANT:
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APPLICANT:
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                                                                                                                                                                                                                                         APPLICANT: La Rosa, Thomas J. APPLICANT: Kovalic, David K. APPLICANT: Zhou, Yihua
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: La Rosa, Thomas APPLICANT: Kovalic, David
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FEATURE:
OTHER INFORMATION: Clone ID: MRT4577_115198C.1.pep
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Oryza sativa
ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                             INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                          279 VGDLGD 284
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                                                                                                                                                                                           Boukharov, Andrey A.
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Pred. No. 1.1e+03;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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APPLICANT: Zhou, Yihuad

APPLICANT: Kovalic, David K.

APPLICANT: Screen, Steven E

APPLICANT: Tabaska, Jack E

APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION UNMBER: US/10/425,114

CURRENT APPLICATION UNMBER: US/10/425,114

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 73128

SEQ ID NO 45513

LENGTH: 422

TYPE: """
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 2002-05-29
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2001-05-30
PRIOR APPLICATION NUMBER: JP 2001-272697
PRIOR FILING DATE: 2001-08-02
NUMBER OF SEQ ID NOS: 15109
SEQ ID NO 14669
LENGTH: 393
Typer: 507
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; OTHER INFORMATION: Clone ID: PAT_MRT4530_100035C.1.pep US-10-437-963-102521
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; OTHER INFORMATION: Clone ID: 700451402_FLI.pep US-10-425-114-45513
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APPLICANT: ISHIKAWA, JUN
APPLICANT: ISHIKAWA, HIROSHI
APPLICANT: SHIBA, TADAYOSHI
APPLICANT: SAKAKI, YOSHIYUKI
APPLICANT: HATTORI, MASAHIRA
APPLICANT: HATTORI, MASAHIRA
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-262
CURRENT APPLICATION UMBER: US/10/156,761
CURRENT FILING DATE: 2002-05-29
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Best Local Similarity 66.7%;
Matches 4; Conservative
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Best Local Similarity 66.7%;
Matches 4; Conservative
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Publication No. US20030119018A1
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APPLICANT: IKEDA, HARUO
APPLICANT: ISHIKAWA, JUN
APPLICANT: HORIKAWA, HIRO
                                     ORGANISM: Zea mays FEATURE:
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Pred. No. 1.2e+03;
2; Mismatches 0; Indels
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Pred. No. 1.4e+03;
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Query Match 90.6%; Score 29; DB 15; Length 422;
Best Local Similarity 66.7%; Pred. No. 1.5e+03;
Matches 4; Conservative 2; Mismatches 0; Indels 0; Gaps
Qy 1 IGDIGD 6
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Db ' 246 LGDVGD 251

Search completed: July 27, 2005, 01:24:19 Job time : 154 secs



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Maximum
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	89	7	თ	_U	4.	ω	N	ַ	i bo
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T23006	T36623	H83610	H81283	AG3239	A70914	E86209	A36046	S74648	S60161	861986	H69143	KIBSFF	AI2674	F70485	H71052	A71663	D97810	C75597	T29528	A42579	JX0259	A31583	A40326	A53812	B35703	S16584	F97008	D36943	ΙĐ
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E83043	C82640	G81785	D97094	G97778	G84722	H69007	F84481	T15917	D69108	F95938	E82523	D69188	T06376	D84634	T36771	F95321	C28379	AE1305	T34919
probable glycosyl	•		probable oxidoredu	cell surface antig	hypothetical prote	tungsten formylmet	Mutator-like trans	hypothetical prote	phycocyanin alpha	probable transcrip	hypothetical prote	nitrogen regulator	ferredoxin-NADP re	hypothetical prote	probable integral	nitrogenase (EC 1.	nitrogenase (EC 1.	conserved hypothet	oxidoreductase - S

ALIGNMENTS

RESULT 1 D36943

A; McLecule type: DNA
A; Molecule type: DNA
A; Molecule type: DNA
A; Molecule type: DNA
A; Cross-references: UNIPROT: Q46589; GB:L26013
A; Cross-references: UNIPROT: Q46589; GB:L26013
A; Cross-references: UNIPROT: Q46589; GB:L26013
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A; Cross-references: UNIPROT: Q46589; GB:L26013
A; Cross-references: UNIPROT: Q46589; Ś chemoreceptor protein dcrD - Desulfovibrio vulgaris (strain Hildenborough) (fragment)
C;Species: Desulfovibrio vulgaris
C;Date: 07-Jul-1995 #sequence_revision 07-Jul-1995 #text_change 09-Jul-2004
C;Accession: D36943 밁 Query Match Best Local 9 Matches N 1 IGDIGD 6 6 Similarity 100.0%; Score 32; Dilarity 100.0%; Pred. No. 9; Conservative 0; Mismatches residue 9 as Ala, GGT fo as His, AAC for residue DB 2; Length 71; 0; Indels GGT for residue 47 as esidue 66 as Ser, GTC 1 0 Gaps 0 for

RESULT 2 F97008

hypothetical protein CAC0881 [imported] - Clostridium acetobutylicum C;Species: Clostridium acetobutylicum C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 09-Jul-2004 C;Accession: F97008

R;Nolling, J.; Breton, G.; Omelchenko, M.V.; Markarova, K.S.; Zeng, Q.; Gibson, R.; Lee.; Daly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R.
J. Bacteriol. 183, 4823-4838, 2001

A;Title: Genome Sequence and Comparative Analysis of the Solvent-Producing Bacterium Clo A;Reference number: A96900; MUID:21359325; PMID:21359325 A;Accession: F97008 A;Status: preliminary

A;Molecule type: DNA A;Residues: 1-190 «KUR» A;Residues: 1-190 «KUR» A;Cross-references: UNIPROT:Q97KN6; GB:AE001437; PIDN:AAK78857.1; PID:g15023777; GSPDB:C A;Experimental source: Clostridium acetobutylicum ATCC824 C;Genetics:

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cofilin, muscle - mouse
C;Species: Nus musculus (house mouse)
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994
C;Date: 08-06-1994 #sequence_revision 07-Oct-1994
C;Accession: A53812
R;Ono, S:; Minami, N.; Abe, H.; Obinata, T.
J. Biol. Chem. 269, 15280-15286, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        coat protein - peanut stunt virus
C;Species: peanut stunt virus
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
C;Accession: S16584
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S16584
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Plant Mol. Biol. 17, 175-177, 1991
A;Title: Nucleotide sequence analysis of a cDNA clone encoding A;Reference number: S16584; MUID:91329730; PMID:1868220
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A; Residues: 1-166 < ABE>
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Biochemistry 29, 7420-7425, 1990
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C;Date: 12-Oct-1990 #sequence_revision 27-Jun-1994 #text_change
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A; Residues: 1-225 <NAI>
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C-reactive protein precursor - mouse C;Species: Mus musculus (house mouse) C;Date: 28-Feb-1990 #sequence_revision C;Accession: A31583; S08286
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C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 09-Jul-2004
C;Accession: A40326; B40326
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A;Molecule type: mRNA
A;Residues: 1-166 <ON
A;Cross-references: UNIPROT:P14847; GB:X13588; R;Whitehead, A.S.; Zahedi, K.; Rits, M.; Morte Biochem. J. 266, 283-290, 1990
A;Title: Mouse C-reactive protein. Generation A;Reference number: S08286; MUID:90179719; PMI
                                                                                                                         R;Ohnishi, S.; Maeda, S.; Nishiguchi, S.; Arao, T.; Shin Biochem. Biophys. Res. Commun. 156, 814-822, 1988 A;Title: Structure of the mouse C-reactive protein gene. A;Reference number: A31583; MUID:89050112; PMID:3190681 A;Accession: A31583
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Biochemistry 30, 9531-9538, 1991
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C;Superfamily: cofilin
C;Keywords: actin binding; muscle
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C;Superfamily: C-reactive
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A;Cross-references: GB:J05343; GB:S56005
C;Comment: This protein, unlike its homolog
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A; Residues: 1-225 < DOW>
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A;Note: the authors translated the codon CTA for residue
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                                                      GB:X13588; NID:g50571; M.; Mortensen, R.F.; l
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A; Introde: --, - C; Superfamily: C-reactive
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A:Molecule type: mRNA
A:Residues: 1-133,'P',135-225 <WHI>
                                                                                                                                                                                                             R;Rassouli, M.; Sambasivam, H.; Azadi, P.; Dell, A.; Morris, H.R. J. Biol. Chem. 267, 2947-2954, 1992
A;Title: Derivation of the amino acid sequence of rat C-reactive A;Reference number: A42579; MUID:92147634; PMID:1737750
A;Accession: A42579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Superfamily: C-reactive protein
C;Keywords: acute phase; liver; pentamer; pentraxin; plasma
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-225/Product: C-reactive protein #status predicted <MET
F;55-116/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Title: Structure, expression, and evolution of guinea A;Reference number: JX0259; MUID:93252746; PMID:8486600 A;Accession: JX0259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C-reactive protein precursor - guinea pig
C;Species: Cavia porcellus (guinea pig)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change
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                                                                                                      A;Note: sequence extracted from NCBI backbone (NCBIP:80758) C;Superfamily: C-reactive protein C;Keywords: acute phase; liver; pentamer; pentraxin; plasma
                                                                                                                                                                                                                                                                                                                    C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                         N;Alternate names: CRP
C;Species: Rattus norvegicus (Norway rat)
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A; Residues: 1-225 < RUB:
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R;Rubio, N.; Sharp, P.M.; Rits, M.; Zahedi, K.; Whitehead, A.S.
J. Biochem. 113, 277-284, 1993
S
                                                                                                                                            A;Cross-references: UNIPROT:P48199; GB:M83176; NID:g203591; A;Note: sequence extracted from NCBI backbone (NCBIP:80758)
                                                                                                                                                                                                                                                                                                                                                             C-reactive protein precursor - rat
N;Alternate names: CRP
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                                                                                                                                                                            A; Residues: 1-230 < RAS>
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R;Wu, X.; Le, T.T. submitted to the EMBL Data Library, November 1996 committed to the sequence of C. elegans cosmid
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DNA topoisomerase (ATP-hydrolysing) (EC 5.99.1.3) - Rickettsia conorii (str: C;Species: Rickettsia conorii (c;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 09-Jul-2004 C;Accession: D97810 C;Accession: D97810 R;Ogata, H.; Audic, S.; Renesto-Audiffren, P.; Fournier, P.E.; Barbe, V.; Science 293, 2093-2098, 2001
                                                                                                                  RESULT 12
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A; Accession: T29528
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C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
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A;Molecule type: DNA
A;Residues: 1-364 <WHI>
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A;Experimental source: strain Bristol N2; clone F27C1
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C; Superfamily: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                          A; Experimental source:
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;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999
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, L.; Utterback,
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Barbe, V.; Samson,

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C;Species:
C;Date: 21-
                                                                                                                                                                                             M.; Ohfuku, Y.; Funahashi, T.; Tanaka, T.; Kuqon, Y.; IDNA Res. 5, 55-76, 1998
A;Title: Complete sequence and gene organization of the A;Reference number: A71000; MUID:98344137; PMID:9679194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Title: The genome sequence of Rickettsia prowazekii and the origin of mitochondria. A;Reference number: A71630; MUID:99039499; PMID:9823893
A;Accession: A71663
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R;Andersson, S.G.B.; Zomorodipour, A.; Andersson, J.O.; Sicheritz-Ponten, T.;
Nature 396, 133-140, 1998
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C;Superfamily: DNA topoisomerase
C;Keywords: isomerase
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                                                                                                                                                                                                                                                           R;Kawarabayasi, Y.; Sawada, M.; M.; Ohfuku, Y.; Funahashi, T.;
                                                                                                                                                                                                                                                                                             C;Date: 14-Aug-1998 #sequence_revision 14-Aug-1998 #text_change 12-Jul-2004 C;Accession: H71052
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A; Residues: 1-807 < KUR>
                                                                                                      A;Cross-references: UNIPROT:058846; GB:AP000005; NID:g3236132; PIDN:BAA30218.1; A;Experimental source: strain OT3
                                                                                                                                      A; Residues: 1-156 < KAW>
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C;Species: Pyrococcus horikoshii
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;Experimental source: strain Madrid E
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J.; Kushida,
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C;Gene
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                                                                                                                                                                                                                                                                                  A; Title: The Genome of the Natural Genetic A; Reference number: AB2577; MUID:21608550; A; Accession: AI2674
                                                                                                                                                                                                                                                                                                                                                                                                           C;Accession: AI2674
R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, erage, G.; Gillet, W.; Grant, C.; Guenthner,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Aquifex aeolicus
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 09-Jul-2004
C;Accession: F70485
                                                                                                                                                                                                           A;Molecule type: DNA
A;Residues: 1-208 <KUR>
A;Cross-references: UNIPROT:Q8UH84; GB:AE008688;
                                                                                                                                                                                                                                                                                                                                                       ; Karp, P.; Romero, P.; Zhang, S.
Science 294, 2317-2323, 2001
A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.; Gordon-Kamm,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hypothetical protein Atu0799 [imported] - Agrobacterium tumefaciens (strain C;Species: Agrobacterium tumefaciens C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 09-Jul-2004
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164
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Eur. J. Biochem. 108, 587-597, 1980
A, Title: Phosphofructokinase: complete amino-acid sequence A, Reference number: A91097; MUID:81003881; PMID:6447595
A, McCession: A00605
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A;Reference number: A93208; MUID:79199719; PMID:156307
A;Contents: annotation; X-ray crystallography, 2.4 angstroms
A;Note: the 6-phosphate group of fructose 6-phosphate (FGP) is bound by His-246 from one d to Asp-124; the sugar ring is bound by Arg-249, Met-166, Glu-219, and Asp-124. ATP is A;Note: the sugar ring is bound by Arg-249, Met-166, Glu-219, and Asp-124. ATP is A;Note: the main interaction with ADP appears to be with the beta-phosphate group, which and Thr-155; the phosphate group of phosphoenolpyruvate appears to bind in the same pl
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Gene 54, 65-71,
                                                                                                                              C;Accession: H69143

R;Smith, D.R.; Doucette-Stamm, L.A.; Deloughery, C.; Lee, H.; Dubois, J.; Aldredge, T.; Qiu, D.; Spadafora, R.; Vicaire, R.; Wang, Y.; Wierzbowski, J.; Gibson, R.; Jiwani, i., Spadafora, Daniels, C.J.; Mao, J.; Rice, P.; Noelling, J.; Reeve, J.N. ki, S.; Church, G.M.; Daniels, C.J.; Mao, J.; Rice, P.; Noelling, J.; Reeve, J.N. J. Bacteriol. 179, 7135-7155, 1997

J. Bacteriol. 179, 7135-7155, 1997

A;Title: Complete genome sequence of Methanobacterium thermoautotrophicum Delta H: fur A;Reference number: A69000; MUID:98037514; PMID:9371463
                                                                                                                                                                                                                                                                                                         coenzyme F420-reducing hydrogenase, beta subunit homolog - Methanobacterium thermoautotrophicum C;Species: Methanobacterium thermoautotrophicum C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Start codon: GTG
C;Superfamily: ATP-dependent phosphofructokinase, prokaryotic type; 6-phosphofructokinase
C;Keywords: allosteric regulation; ATP; glycolysis; homotetramer; phosphotransferase
F;4-278/Domain: 6-phosphofructokinase 1 homology <6PF1>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Sakai, H.; Ohta, T.
Eur. J. Blochem. 211, 851-859, 1993
Eur. J. Blochem. 211, 851-859, 1993
A;Title: Molecular cloning and nucleotide sequence of the gene sphofructokinase and pyruvate kinase constitute an operon.
A;Reference number: S29783; MUID:93170322; PMID:8436141
A;Accession: S29784
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A;Residues: 1-11,'N',13-35,39-42,'V',44-81,'E',83-94,'Q',96-224,'L',226-263,'SA',266-31
A;Note: this enzyme is regulated by ADP and phosphoenolpyruvate, in contrast to the euk
R;Evans, P.R.; Hudson, P.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: DNA
A; Residues: 1-319 <FRE>
A; Cross-references: UNIPROT: P00512; GB: M15643; NID: g143311; PIDN: AAA22656.1;
R; Kolb, E.; Hudson, P.J.; Harris, J.I.
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A; Residues: 300-319
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                              A; Molecule type: DNA
A; Residues: 1-406 < MTH>
                                                                                                        A; Reference number: A; Accession: H69143
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                                                                               A; Status: preliminary; nucleic
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A;Cross-references: UNIPROT:O26441; GB:AE000819; GB:AE000666;
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;Date: 30-Sep-1980 #sequence_revision 14-Jul-1994 #text_change 09-Jul-2004
;Accession: A27474; A00605; S29784
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NID:g2621396; PIDN:AAB8484
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  transcription factor relB - African clawed frog
C;Species: Xenopus laevis (African clawed frog)
C;Date: 15-Feb-1996 #sequence_revision 23-Feb-1;
C;Accession: S60161; S60162
R;Suzuki, K.; Yamamoto, T.; Inoue, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Sterky, F.; Holmberg, A
Yeast 12, 1091-1095, 1996
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A; Residues: 1-478 < STE>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Accession: S61986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N; Alternate names:
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A;Cross-references: SGD:S00055
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A;ACCEBBIONA
A;Molecule type; DNA
A;Rosidues: 1-478 <PET>
A;Rosidues: 1-478 <PET>
A;Cross-references: EMBL:Z74911; NID:g1420091; PIDN:CAA99191.1; PID:g1420092;
A;Cross-references: ExpEt:Z74911; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:g1420091; PID:
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A;Gene: MTH341
C;Superfamily: formate dehydro
                                                                                                                                                                                C;Keywords: hydrolase; serine proteinase F;1-20/Domain: signal sequence #status predicted <SIG>F;21-478/Product: subtilisin-like proteinase YSP3 #status predicted F;204-421/Domain: subtilisin homology <SBT>F;203-421/Domain: subtilisin homology <SBT>F;213,245,407/Active site: Asp, His, Ser #status predicted
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A;Cross-references: EMBL:U43491; NID:g1150992; PI
A;Note: the nucleotide sequence was submitted to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Title: The sequence of a 30 kb fragment on A; Reference number: 872130; MUID: 97051599; P
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C; Superfamily: subtilisin; subtilisin homology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SGD:S0005529; MIPS:YOR003w
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Pred. No.
                                    Score 30;
Pred. No. 1
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#text_change 09-Jul-2004

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C;Date: 16-Nov-1990 #sequence_revision 13-Sep-1991 #text C;Accession: A36046 R;Exposito, J:Y:; Garrone, R. Pxposito, J:Y:; Garrone, R. Pxposito, J:Y:; Garrone, R. Proc. Natl. Acad. Sci. U.S.A. 87, 6669-6673, 1990 A;Title: Characterization of a fibrillar collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification of the collagen gene identification gene identification gene identification gene identification gene identification gene identification gene identification gene identification gene identification gene identification gene identification gene identification gene identification gene identification gene identification general gene identification general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general general
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Nucleic Acids Res. 23, 4664-4669, 1995
A;Title: Molecular cloning of cDNA encoding the Xenopus
A;Reference number: S60162; MUID:96103597; PMID:8524658
A;Accession: S60162
A;Status: nucleic acid sequence not shown
A;Molecule type: mRNA
A;Residues: 78-367 <SUW>
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C;Superfamily: 4-alpha-glucanotransferase
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A;Residues: 1-497 <SUZ>
A;Cross_references: UNIPROT:P51510;
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                                                                                                                                                    collagen alpha chain - freshwater sponge (Ephydatia muelleri) (fragment) (;Species: Ephydatia muelleri C;Decies: Ephydatia muelleri C;Date: 16-Nov-1990 #sequence_revision 13-Sep-1991 #text_change 09-Jul-2 C;Accession: A36046
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C;Genetics:
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A;Residues: 1-505 <KAN>
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K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamada, M.; Yasud
A Res. 3, 109-136, 1996
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Pred. No. 2e+0
1; Mismatches
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Pred. No. 2.1e+02;
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submitted to
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       A;Experimental source: strain H37Rv C;Genetics:
                                                                        A;Cross-references: UNIPROT:P71714; GB:Z80775; GB:AL123456; NID:g3250715; PIDN:CAB02535.1
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A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W. A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis. A;Reference number: A86141; MUID:21016719; PMID:11130712
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Nature 408, 816-820, 2000
A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, B.; Kim, C., A;Authors: J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Maiti, R.; Marziali, C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Marziali, G.A.; Li, J.H.; Li, Rowley, D.; Sakano, H.
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
                                                                           A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete A;Reference number: A70500; MUID:98295987; PMID:9634230
A;Accession: A70914
                                                                                                                                                                           R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, 537-544, 1998
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A; Residues: 1-84 <STO>
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A; Residues: 1-547 < EXP>
A; Molecule type: DNA
A; Residues: 1-173 < COL>
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C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 09-Jul-2004
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                                                A;Status: preliminary; nucleic acid sequence not
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                                                                                                                                                                                                                           , D.; Gorαομ,
N.; Holroyd,
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Marziali,
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C;Date: 11-Jan-2002 #sequence revision 11-Jan-2002 #text_change 09-Jul-2004 C;Accession: AG3239  
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C;Superfamily:
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A;Residues: 1-218 <PAR>
A;Cross-references: UNIPROT:O9PMR3; GB:AL139078; GB:AL111168; NID:g6968723; PIDN:CAB7381
A;Exmerimental source: serotype O2, strain NCTC 11168
                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Title: The genome sequence of the food-borne pathogen Campylobacter jejuni reveals hypa;Reference number: A81250; MUID:20150912; PMID:10688204
A;Accession: H81283
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A;Title: The Genome of the Natural Genetic A;Reference number: AB2577; MUID:21608550; A;Accession: AG3239
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Karp, P.; I
Science 294,
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A;Cross-references: UNIPROT:Q8U638; GB:AE008690; PIDN:AAL46333.1; PID:g17744120; GSPDB:
A:Experimental source: strain C58 (Dupont)
                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                     A;Status: preliminary
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1.3e+02;
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T.; Levy, R.; Li, M.; McClel
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                             A; Molecule type: DNA
A; Residues: 1-338 <WIL>
A;Cross-references: UNIPROT:Q21039; EMBL:Z68005; PIDN:CAA91992.1; GSPDB:GN00028; CESP:F5
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C;Accession: H83610
R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.;
R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.;
adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, R
adman, S.; Olson, M.V.
i, Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
                          C;Date: 15-00-10
C;Accession: T23006
R;Kershaw, J.
Rubmitted to the EMBL Data Library,
submitted to the EMBL Data Library,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, A;Reference number: A82950; MUID:20437337; PMID:10984043 A;Accession: H83610
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H83610
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C;Species: Streptomyces coelicolor
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004
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C;Superfamily: argina
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C;Date: 15-Sep-2000 #sequence_revision
                                                                                                             hypothetical protein F59F3.2 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
                                                                                                                                                                            RESULT
T23006
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A; Residues: 1-320 <OLI>
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A;Accession: T36623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Oliver, K.; Harris, D.; Bentley, S.D.; submitted to the EMBL Data Library, June
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Accession: T36623
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A; Residues: 1-318 <STO>
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                      A; Reference number:
A; Accession: T23006
A;Status: preliminary; translated from GB/EMBL/DDB:
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Pred. No. 2e+(
2; Mismatches
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                                                                                                                                                                                                                                                                                                                                     Score 29; DB 2;
Pred. No. 2e+02;
2; Mismatches
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                                                               November 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL:AL078610; PIDN:CAB44391.1;
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1999
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2e+02;
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K.R.; Kas, A.; Larbig, K.; Lim,
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Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J. A;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
                                                                                                                                                                                                                                                                                                                                                                                                                            conserved hypothetical proteins homolog lmol845 [imported] - Listeria monocytogenes (str C;Species: Listeria monocytogenes C;Species: Listeria monocytogenes C;Date: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change 09-Jul-2004 C;Accession: AE1305 C;Accession: AE1305 R;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.; Baquero, F.; Berche, P.; Bloecker, Dominguez-Bernal, G.; Duchaud, E.; Durand, L.; Dussurget, O.; Entian, K.D.; Fsihi, H.
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C;Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text_change 09-Jul-:
C;Accession: T34919
R;Oliver, K.; Harris, D.; Parkhill, J.; Barrell, B.G.; Rajandream, M.A.
submitted to the EMBL Data Library, January 1998
A;Reference number: Z21558
A;Accession: T34919
A;Status: preliminary; translated from GB/EMBL/DDBJ
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A;Map position: X
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                                                                                                                                                                                              A;Cross-references: UNIPROT:Q8Y655; GB:NC_003210; PIDN:CAC99923.1; PID:g16411299; GSPDB
A;Experimental source: strain EGD-e
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C;Superfamily: Escherichia coli UDPglucose 4-epimerase; UDPglucose
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A; Residues: 1-347 <OLI>
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                                                                                                                                                                                                                                    A;Residues: 1-427 <GLA>
                                                                                                                                                                                                                                                                          A;Status: preliminary
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Best Local S
Matches 4
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Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text_change 09-Jul-2004
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                                                                          Score 29; DB 2;
Pred. No. 2.7e+02;
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Pred. No. 2.2e+02;
2; Mismatches (
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Pred. No. 2.1e+02;
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RESULT
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A;Gene: fixF
C;Superfamil:
C;Keywords: i
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A;Authors: Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, hebault, P.; Vandenbol, M.; Vorholter, F.J.; Weidner, S.; Wells, D.H.; Wong, K.; Yeh, K.(A;Title: The composite genome of the legume symbiont Sinorhizobium meliloti.
A;Reference number: A96039; MUID:21368234; PMID:11474104
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C,Species: Rhizobium meliloti
C,Date: 30-Jun-1989 #sermenca ------'
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F95321
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A;Title: Rhizoblum meliloti nifN (fixF) gene is part of
A;Reference number: A91851; MUID:88058743; PMID:3316182
                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-441 <KUR>
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C;Species: Sinorhizobium meliloti
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A; Residues: 1-441 < AGU>
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Accession: C28379
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Pred. No.
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C;Accessic
R;Seeger,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypothetical protein At2g24250 [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Peb-2001 #sequence_revision 02-Peb-2001 #text_change 09-Jul-2004
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A;Residues: 1-684 <SEE>
A;Cross-references: UNIPROT:Q9S282;
A;Experimental source: strain A3(2)
C;Genetics:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probable integral membrane protein - Streptomyces coelicolor
C;Species: Streptomyces coelicolor
C;Date: 03-Dec_1999 #sequence_revision 03-Dec-1999 #text_change
                                                                                                                                                   R;Marraffini, L.A.; Ceccarelli, E.A. submitted to the EMBL Data Library, April 1 A;Description: Involvement of tyrosine-114
                                                                                                                                                                                                      C;Species: Pisum sativum (garden pea)
C;Date: 30-Apr-1999 #sequence_revision
C;Accession: T06376
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A;Accession: T36771
A,Status: preliminary; translated A,Molecule type: mRNA A,Residues: 1-35 <MAR> A,Cross-references: UNIPROT:O65208 C,Genetics: A,Genome: nuclear
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                                                                                                             A; Reference number: Z15636
A; Accession: T06376
                                                                                                                                                                                                                                                                                                      RESULT 36
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A; Residues: 1-695 <STO
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                                                                                                                                                                                                                                                                ferredoxin-NADP reductase (EC 1.18.1.2) -
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Best Local S
Matches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: UNIPROT:Q9ZUHO; GB:AE002093; NID:g4115385; PIDN:AAD03386.1; GSPDB:GN
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                                       UNIPROT: 065208; EMBL: AF057182; NID: g3088653; PIDN: AAC14746.1;
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C;Superfamily: ferredoxin-NADP+ reductase; cytochrome-b5 reductase homology C;Keywords: chloroplast; electron transfer; FAD; flavoprotein; NADP; oxidoreductase;
A;Molecule type: DNA
A;Residues: 1-147 <SIM>A;Residues: 1-147 <SIM>A;Residues: 1-147 <SIM>A;Cross-references: UNIPROT;O9P9Y8; GB:AE004079; GB:AE003849; NID:g9107960; PIDN:AAF8551
A;Cross-references: UNIPROT;O9P9Y8; GB:AE004079; GB:AE003849; NID:g9107960; PIDN:AAF8551
A;Experimental source: strain 9a5c
R;Simpson, A.J.G.; Resinach, F.C.; Arruda, P.; Abreu, F.A.; Acencio, M.; Alvarenga, R.; A
Briones, M.R.S.; Bueno, M.R.P.; Camargo, A.A.; Camargo, L.E.A.; Carraro, D.M.; Carrer, H
as-Neto, B.; Docena, C.; Bl-Dorry, H.; Facincani, A.P.; Ferreira, A.J.S.
as-Neto, B.; Docena, C.; Bl-Dorry, H.; Facincani, A.P.; Ferreira, A.J.S.
as-Neto, B.; Docena, C.; Bl-Dorry, H.; Fraga, J.S.; Franca, S.C.; Franco, M.C.; Frohm
A;Authbors: Ferreira, V.C.A.; Ferro, J.A.; Fraga, J.S.; Franca, S.C.; Franco, A.G.; Frohm
A;D.; Junqueira, M.L.; Kemper, B.L.; Kitajima, J.P.; Krieger, J.B.; Kuramae, B.E.; Laigr
Chado, M.A.; Macdeira, A.M.B.N.; Madeira, H.M.F.; Marrino, C.L.; Marques, M.V.; Martins, E.M.F.; Matsukuma, A.Y.; Menck, C.F.M.; Miracca, B.C.; Miyaki, C.Y.,
A;Authors: Martins, E.M.F.; Matsukuma, A.Y.; Menck, C.F.M.; Miracca, B.C.; Palmieri, D.A
,F.G.; Nunes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, R.C.; Palmieri, D.A
                                                                                                                                                                                                                                                                                                                                                                                                                R;anonymous, The Xylella fastidiosa Consortium of the Organization for Nature 406, 151-157, 2000
A;Title: The genome sequence of the plant pathogen Xylella fastidiosa. A;Reference number: A82515; MUID:20365711; PMID:10910347
A;Note: for a complete list of authors see reference number A59328 bel
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C;Keywords: phospho
F;54/Binding site:
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C;Date: 05-Dec-1997 #sequence_revision
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hypothetical protein XF2732 [imported] - Xylella fastidiosa (strain 9a5c)
C;Species: Xylella fastidiosa
C;Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 09-Jul-20
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;Superfamily: regulatory protein P-II
;;Superfamily: regulatory protein P-II
;;Keywords: phosphoprotein; signal transduction
;;Keywords: phosphoprotein | signal transduction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ,Smith, D.R.; Doucette-Stamm, L.A.; Deloughery, C.; Lee, H.; Dubois, J.; Aldredge, T. Qiu, D.; Spadafora, R.; Vicaire, R.; Wang, Y.; Wierzbowski, J.; Gibson, R.; Jiwani, i, S.; Church, G.M.; Daniels, C.J.; Mao, J.; Rice, P.; Noelling, J.; Reeve, J.N. Bacteriol. 179, 7135-7155, 1997
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Matches
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                               see reference number A59328 below
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le+02;
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R;Smith, D.R.; Doucette-Stamm, L.A.; Deloughery, C.; Lee, H.; Dubois, J.; Aldredge, T.; Qiu, D.; Spadafora, R.; Vicaire, R.; Wang, Y.; Wierzbowski, J.; Gibson, R.; Jiwani, ki, S.; Church, G.M.; Daniels, C.J.; Mao, J.; Rice, P.; Noelling, J.; Reeve, J.N. J. Bacteriol. 179, 7135-7155, 1997
A;Title: Complete genome sequence of Methanobacterium thermoautotrophicum Delta H: fur A; Reference number: A69000; MUID: 98037514; PMID: 9371463
A; Accession: D69108
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C;Accession: F95938
                               A;Status: preliminary; nucleic acid sequence A;Molecule type: DNA A;Residues; 1-183 <MTH>
                                                                                                                                                                                                       phycocyanin alpha phycocyanobilin lyase CpcE - Methanobacterium thermoautotrophicum C;Species: Methanobacterium thermoautotrophicum C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004 C;Accession: D69108
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A; Residues: 1-154 < KUR>
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A;Authors: da Silva, A.C.R.; da Silva, F.R.; da Silva, A.M.; Silva Jr., W.A.; da Silveix
M.; Tsuhako, M.H.; Vallada, H.; Van Sluys, M.A.; Verjovski-Almeida, S.; Vettore, A.L.; Z
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                ;Cross-references:
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Pred. No. 1.3e+02;
2; Mismatches 0
 GB:AE000934; GB:AE000666; NID:g2622924; PIDN:AAB862:
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Bubmitted to the EMBL Data Library, July 1995

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C;Superfamily: glycine-rich RNA-binding protein; ribonucleoprotein repeat homology F;20-87/Domain: ribonucleoprotein repeat homology <RRM>
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A;Cross-references: UNIPROT:Q09511; EMBL:U23484; NID:g733597; PID:g733604; PIDN:AAC46767
A;Experimental source: strain Bristol N2
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C;Species: Caenorhabditis elegans
C;Species: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                               A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana A;Reference number: A84420; MUID:20083487; PMID:10617197 A;Accession: F84481 A;Accession: F84481 A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                     Mutator-like transposase [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 09-Jul-2004
C;Accession: F84481
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A;Gene: MTH1806
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A;Map position: 2
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A; Residues: 1-212 <STO>
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                                                                                                                                                                                                                                                                                                                                                 Nature 402, 761-768, 1999
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C;Species: Ar
C;Date: 02-Fel
R;Ogata, H.; Audic, S.; Renesto-Audiffren, P.; Fournier, P. science 293, 2093-2098, 2001
A;Title: Mechanisms of Evolution in Rickettsia conorii and A;Reference number: A97700; MUID:21442074; PMID:11557893
A;Accession: G97778
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A;Cross-references: UNIPROT:026209; GB:AE000800; GB:AE000666; NID:g2621130; PIDN:AAB8460
A;Experimental source: strain Delta H
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                                                                                         cell surface antigen homolog [imported] - Rickettsia conorii (strain Malish C;Species: Rickettsia conorii (C;Species: Rickettsia conorii (C;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 09-Jul-2004 C;Accession: G97778
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A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana.
A;Reference number: A84420; MUID:20083487; PMID:10617197
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                                                                                                                                                                                                                                                                                                                                                                                                    Superfamily:
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Gene: At2g31600
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genetics
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                                                                                                                                                                                                                                                                                                                          2.2e+02;
hes 0;
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2.1e+02;
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R;Nolling, J.; Breton, G.; Omelchenko, M.V.; Markarova, K.S.; Zeng, Q.; Gibson, J.; Daly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R.
J. Bacteriol. 183, 4823-4838, 2001
A;Title: Genome Sequence and Comparative Analysis of the Solvent-Producing Bacte A;Reference number: A96900; MUID:21359325; PMID:21359325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Molecule type: DNA
A;Residues: 1-256 <KUR>
A;Cross-references: UNIPROT:Q92HY9; GB:AE006914; PIDN:AAL03169.1; PID:g15619717; GSPDB:G
                                                                                                                                                                                                                                                                   R;Parkhill, J.; Achtman, M.; James, ; Holroyd, S.; Jagels, K.; Leather, Nature 404, 502-506, 2000
                                                                                                                                                                                                                                                                                                                      hypothetical protein NMA2138 [imported] - Neisseria meningitidis (strain Z2491 C;Species: Neisseria meningitidis C;Date: 05-May-2000 #sequence_revision 05-May-2000 #text_change 09-Jul-2004 C;Accession: G81785
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C; Superfamily: ribitol dehydrogenase; short-chain alcohol dehydrogenase homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 09-Jul-2004 C;Accession: D97094
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probable oxidoreductase, ortholog of GSP39 B. subtilis CAC1576 [imported] - Clostridium
C;Species: Clostridium acetobutylicum
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                                                                                              A; Experimental source: C; Genetics:
                                                                                                                                                                                                                            A; Title: Complete DNA sequence of a serogroup A strain of A; Reference number: A81775; MUID:20222556; PMID:10761919
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                                                                                                                                                   A;Molecule type: DNA
A;Residues: 1-295 <PAR>
                                                                                                                                                                                                        A; Accession: G81785
                                                                                                                                                                                        A;Status: preliminary
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;Experimental source: Clostridium a
                                                                                                                Cross-references: UNIPROT:Q9JST9;
Experimental source: serogroup A,
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5; Conserv
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     Conservative
100.0%; **
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100.0%; Pred. No. 2.
tive 0; Mismatches
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     Score 28; DB; Pred. No. 2.9
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                                                                                                                    GB:AL162758;
strain Z2491
                                                                                                                                                                                                                                                                                     K.D.; Bentley,
S.; Moule, S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 2; L. 2.8e+02;
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2.5e+02;
                           2 B
                         B 2; L
                                                                                                                                     GB:AL157959; NID:g7380672; PIDN:CAB8535
                                                                                                                                                                                                                                                                                       S.D.; Churcher, C.;
Mungall, K.; Quail,
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                                           Length
         Indels
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M.A.;
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Rajandream,
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A; Molecule type: DNA
A; Residues: 1-296 <SIM>
A; Cross-reference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Experimental source: strain 9a5c

R;Simpson, A.J.G.; Reinach, F.C.; Arruda, P.; Abreu, F.A.; Acencio, M.; Alvarenga, R.; A
Briones, M.R.S.; Bueno, M.R.P.; Camargo, A.A.; Camargo, L.E.A.; Carraro, D.M.; Carrer, H
Briones, Docena, C.; El-Dorry, H.; Facincani, A.P.; Ferreira, A.J.S.

as-Neto, E.; Docena, C.; El-Dorry, H.; Facincani, A.P.; Ferreira, A.J.S.

submitted to GenBank, June 2000

A;Authors: Ferreira, V.C.A.; Ferro, J.A.; Fraga, J.S.; Franca, S.C.; Franco, M.C.; Frohm
J.D.; Junqueira, M.L.; Kemper, E.L.; Kitajima, J.P.; Krieger, J.E.; Kuramae, E.E.; Laigr
Chado, M.A.; Madeira, A.M.B.N.; Madeira, H.M.F.; Maxino, C.L.; Marques, M.V.; Marrins, E
A;Authors: Martins, E.M.F.; Matsukuma, A.Y.; Menck, C.F.M.; Miracca, E.C.; Miyaki, C.Y.;

F.G.; Nunes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, R.C.; Palmieri, D.A
Rodrigues, V.; Rosa, A.J. de M.; de Rosa Jr., V.E.; de Sa, R.G.; Santelli, R.V.; Sawasak
A;Authors: da Silva, A.C.R.; da Silva, F.R.; da Silva, A.M.; Silva Jr., W.A.; da Silveira
M.; Tsuhako, M.H.; Vallada, H.; Van Sluys, M.A.; Verjovski-Almeida, S.; Vettore, A.L.; Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hypothetical protein XF1770 [imported] - Xylella fastidiosa (strain 9a5c)
C;Species: Xylella fastidiosa
C;Date: 18-Nag-2000 #sequence_revision 20-Aug-2000 #text_change 09-Jul-2004
C;Accession: C82640
C;Accession: C82640
R;anonymous, The Xylella fastidiosa Consortium of the Organization for Nucleotide Sequent Nature 406, 151-157, 2000
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A; Status: prelimina
                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: DNA
A; Residues: 1-303 <STO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana A;Reference number: A84420; MUID:20083487; PMID:10617197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               M.; Koo, H.;
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Best Local (
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                                                                                                                                                                                                                                                                       Gene: At2g33330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ., X.; Kaul, S.; Rounsley, S.D.; Shea, T.P.; Benito, M.I.; Town, C.D.; Fujii, C.Y.; Oo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon, L. D.; Nierman, W.C.; White, O.; Eisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter,
                                                                                                                 Local Similarity
nes 5; Conserv
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66.7%;
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100.0%; Pr
                                                                                                                                               Score 28; pred. No.
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Nature 406, 959-964, 2000
Nature 406, 959-964, 2000
A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic pathog
A:Reference number: A82950; MUID:20437337; PMID:10984043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probable glycosyl transferase PA4819 [imported] - Pseudomonas aeruginosa (strain C;Species: Pseudomonas aeruginosa C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-Jul-2004 C;Accession: E83043
Search completed: July 27, 2005, 01:21:13 Job time: 42 secs
                                                                                                                                                                                                                                                                                                                                                                                       A;Status: preliminary
A;Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Reference number: A82950; A; Accession: E83043
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A;Experimental source: strain PAO1
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01-NOV-1996 (TrEMBLrel. 01, I
01-OCT-2003 (TrEMBLrel. 25, I
Chemoreceptor (Fragment).
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Desulfovibrio vulgaris.
Bacteria; Proteobacteria; Deltaproteobacteria; Desulfovibrionales;
Desulfovibrionaceae; Desulfovibrio.
NCBI_TaxID=881;
                                                                                                                                                                                                                                                             Deckers H.M., Voordouw G.;
"Identification of a large family of genes for putative chemoreceptor proteins in an ordered library of the Desulfovibrio vulgaris Hildenborough genome.";
J. Bacteriol. 176:351-358(1994).
EMBL; L26013; AAA235881; -..
EMBL; L26013; AAA235881; -..
                                                                                                                                            Pfam; PF00015; MCPsignal; 1.
PRINTS; PR00260; CHEMTRNSDUCR.
PROSITE; PS50111; CHEMOTAXIS_TRANSDUC_2;
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GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0006935; P:chemotaxis; IEA.
GO; GO:0007165; P:signal transduction; IEA.
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InterPro; IPR004090; Me_chemotaxis.
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MEDLINE=21359325; PubMed=11466286;
DOI=10.1128/JB.183.16.4823-4838.2001;
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Q97KN6;
                                                                                                   EMBL; U15730; AAA70087.1;
EMBL; X56544; CAA39885.1;
PIR; S16584; S16584.
                                                                                                                                                                                                       Naidu R.A., Collins G.B., Ghabrial S.A.;
"Nucleotide sequence analysis of a cDNA clone encoding the protein gene of peanut stunt virus.";
Plant Mol. Biol. 17:175-177(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            homology.";
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InterPro; IPR009061; Putativ DNA bind.
Complete proteome; Hypothetical protein.
SEQUENCE 190 AA; 22079 MW; 7D6D3F4A38C36F10 CRC64;
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Clostridium acetobutylicum
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01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Naidu R.A., Hu C.-C.,
"Differentitation of ocucumovirus based on a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Name=coat protein;
Peanut stunt virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Coat protein.
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                            GO:0019028; C:viral capsid; IEA.
GO:0005198; F:structural molecule activity; IEA
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Cucumovirus_coat
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Pred. No. 2e+02;
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Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
Barnay S., Blanchin S., Beckerich J.M., Beyne E., Bleykasten C.,
Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
Bespons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
Nicaud J.M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
Pellenz S., Potier S., Richard G.F., Straub M.L., Suleau A.,
Swennene D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
Zeniou-Meyer M., Zivanovic I., Bolotin-Fukuhara M., Thierry A.,
Boucher C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
Weissenbach J.,
                                                                                                                                                                                                                         Glycosyltransferase;
SEQUENCE 229 AA; ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases EMBL; CR382131; CAG79790.1; -. EMBL; GO:0004743; F:pyruvate kinase activity; IEA. GO: GO:0016757; F:transferase activity, transferring gl: GO: GO:0006096; P:glycolysis; IEA. GO: GO:0009116; P:nucleoside metabolism; IEA.
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SEQUENCE 225 AA;
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PRINTS; PR00222; CUCUMOCOAT.
ProDom; PD001284; Cucumovirus_coat; 1.
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                                                                                                                                                                                                                                                                                                            ProDom;
                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR000836; PRtransferase.
InterPro; IPR001697; Pyruvate_kinas
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"Genome evolution in yeas
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                                                                                   Similarity 6; Conserv
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Thomson N.R., Kieser H., Farvin S., Chandra G., Chen C.W., Collins M., Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth S., Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth S., Fusher C.-H., Kieser T., Larke L., Murphy L.D., Oliver K., O'Neil S., Rabbinowitsch E., Rajandream M.A., Rutherford K.M., Rutter S., Rabbinowitsch E., Rajandream M.A., Rutherford K.M., Rutter S., Taylor K., Seger K., Saunders D., Sharp S., Squares R., Squares S., Taylor K., Seger K., Saunders D., Sharp S., Squares R., Squares S., Taylor K., Seger K., Saunders D., Sharp S., Squares R., Squares S., Taylor K., Seger K., Saunders D., Woodward J.R., Barrell B.G., Parkhill J.,
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Submitted (SEP-2003) to the )
EMBL; AE017227; ALS02563.1;
InterPro; IPR005142; eRF1_3.
Pfam; PF03465; eRF1_3; 1.
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Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
Corynabacterineae; Mycobacteriaceae; Mycobacterium.
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Q9KYG6;
01-OCT-2000 (TrEMBLrel. 15,
01-JUN-2003 (TrEMBLrel. 24,
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                                                                                                                                      SEQUENCE
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Bacteria; Actinobacteridae; Actinomycetales; Streptomycineae; Streptomycetaceae; Streptomyces.
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GO:0016740; F:transferase activity;
GO:0009058; P:biosynthesis; IEA.
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EMBL/GenBank/DDBJ databases
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                                4.7e+02;
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RI Proc. Natl. Acad. Sci. U.S.A. 99:14422-14427(2002).

CC -|- SIMILARITY: Contains 1 histidine kinase domain.

CR EMBL; AE014650; AA024157.1; --

DR GO; GO:0016520; C:membrane; IEA.

GO; GO:0001524; F:ATP binding; IEA.

GO; GO:0001524; F:ATP binding; IEA.

GO; GO:000155; F:two-component sensor molecule activity; IEA.

DR GO; GO:000155; F:two-component sensor molecule activity; IEA.

DR GO; GO:000155; F:two-component sensor molecule activity; IEA.

DR GO; GO:000155; F:two-component sensor molecule activity; IEA.

DR InterPro; IPR003594; ANTPbind_ATPase.

InterPro; IPR003594; ANTPbind_ATPase.

InterPro; IPR003594; His Kinase.

DR InterPro; IPR003661; His Kinase.

DR InterPro; IPR003661; His Kinase.

DR InterPro; IPR003661; His Kinase.

DR InterPro; IPR003661; His Kinase.
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Matches 6
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Pfam; PP00512; HisKA; I.
PRINTS; PR00344; BCTRLSENSOR.
SMART; SM00387; HATPase C; 1.
SMART; SM00388; HisKA; I.
PROSITE; PS50109; HIS KIN; 1.
Complete proteome; Kinase; Phos
                                                                                                                                                                       Q9HKD4 PRELIMINARY;
Q9HKD4;
Q1-MAR-2001 (TrEMBLrel. 16,
Q1-MAR-2001 (TrEMBLrel. 16,
Q1-MAR-2004 (TrEMBLrel. 26,
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Histidine kinase sensor of two-component system.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=216816;
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Bifidobacteriaceae, Bifidobacterium.
Archaeā; Euryarchaeota; Thermoplasmata; Thermoplasmatales; Thermoplasmataceae; Thermoplasma. NCBI_TaxID=2303;
                                                                                                                         Acyl-CoA synthetase related
OrderedLocusNames=Ta0667;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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100.0%; Pred. No. 4.8e+02;
tive 0; Mismatches 0;

    Phosphorylation; Sensory transduction;

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SEQUENCE
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EMBL; AF307843; AAG29840.1; -.

EMBL; AF307842; AAG29839.1; -.

HSSP; 087172; 1ESW.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE-22631581; PubMed-12746519; DOI=10.1104/pp.102.016527; Wattebled F., Ral J.P., Dauvillee D., Myers A.M., James M.G., Schlichting R., Giersch C., Ball S.G., D'Hulst C.; Carrier Chlamydomonas reinhardtii locus required for normal starch granule biogenesis, encodes disproportionating enzyme. Further evidence for a function of alpha-1,4 glucanotransferases during starch granule biographess in green algae.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chlamydomonas reinhardtii.
Eukaryota; Viridiplantae; Chlorophyta; Chlorophyceae; Volvocales;
Chlamydomonadaceae; Chlamydomonas.
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GO; GO:0003824; F:catalytic activity;
GO; GO:0008152; P:metabolism; IEA.
InterPro; IPR000873; AMP-bind.
Pfam; PP00501; AMP-binding; 1.
PRINTS; PR00154; AMPBINDING.
PROSITE; PS00455; AMP_BINDING; 1.
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SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
STRAIN=DSM 1728; PubMed=11029001; DOI=10.1038/35035069;
MEDLINE=20479972; PubMed=11029001; DOI=10.1038/35035069;
                                                                                                                                                                                                                                                  InterPro; IPR003385; Glyco_hydro_7
Pfam; PF02446; 4a_glucanotrans; 1.
TIGRFAMs; TIGR00217; malQ; 1.
                                                                                                                                                                                                                                                                                                                       GO; GO:0004134; F:4-alpha-glucanotransferase activity; IEA.
GO; GO:0016740; F:transferase activity; IEA.
GO; GO:0005975; P:carbohydrate metabolism; IEA.
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-!- SIMILARITY: Belongs to the ATP-dependent AMP-binding
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                                                                                               Similarity 6; Conserv
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Mismatches 0;
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Best Local S
Matches 6
                                       EMBL, ABO04828, BAB95564.1;

GO; GO:0009986; C:cell surface; IEA.
GO; GO:0016020; C:membrane; IEA.
InterPro; IPR011439; DUF1542.
InterPro; IPR011439; FIVAR.
InterPro; IPR011490; FIVAR.
InterPro; IPR005877; Gpos YSIRK.
InterPro; IPR001899; Gram pos_anchor.
Pfam; PF07564; FIVAR; 1.
Pfam; PF07564; FIVAR; 1.
TIGRPAMS; TIGR01167; LPXTG anchor; 1.
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Q87MP0;
01-JUN-2003
01-JUN-2003
01-CCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=22040717; PubMed=12044378; DOI=10.1
Baba T., Takeuchi F., Kuroda M., Yuzawa H.,
Nagai Y., Tawma N., Asano K., Naimi T., Kur
Yamamoto K., Hiramatsu K.;
"Genome and virulence determinants of high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lancet 361:743-749(2003).
EMBL, AP005080; BAC60454.1; -.
InterPro; IPR008941; TPR-like.
Complete proteome; Hypothetical
SEQUENCE 1380 AA; 152808 MW;
                                                                                                                                                                                                                                                                                                                                                   acquired MRSA.";
Lancet 359:1819-1827(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Staphylococcus aureus Bacteria; Firmicutes;
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01-OCT-2002
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STRAIN=RIMD 221063 / Serotype 03:K6;

STRAIN=RIMD 221063 / Serotype 03:K6;

MEDLINE=22508454; PubMed=12620739; DOI=10.1016/S0140-6736(03)12659-1;

Makino K., Oshima K., Kurokawa K., Yokoyama K., Udta T., Tagomori K.,

Iijima Y., Najima M., Nakano M., Yamashita A., Kubota Y., Kimura S.,

Yasunaga T., Honda T., Shinaqawa H., Hattori M., Iida T.;

Yasunaga T., Honda T., Shinaqawa H., Hattori M., Iida T.;

"Genome sequence of Vibrio parahaemolyticus: a pathogenic mechanism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Proteobacteria;
Vibrionaceae; Vibrio.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    distinct from that of V. cholerae.";
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                         TIGRFAMS; TIGRO1167;
TIGRFAMS; TIGRO1168;
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proteome.
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                            YSIRK_signal;
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Bacillales; Staphylococcus
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52197DEF11AF4E7C CRC64;
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uzawa H., Aoki K.-I., Oguchi A.,
i T., Kuroda H., Cui L.,
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RESULT 13
COP2 LOWAZ
ID COP2
AC Q9Y2
AC Q9Y2
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DT 30-M
DT 25-O
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Best Local :
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InterPro; IPR005877; Gpos_YSIRK.
InterPro; IPR001899; Gram_pos_anchor.
Pfam; PF07564; DUF1542; 17.
Pfam; PF04550; YSIRK signal; 1.
TIGREAMs; TIGR01167; LPXTG_anchor; 1.
TIGREAMs; TIGR01168; YSIRK_signal; 1.
Complete proteome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q6G8H1;
05-JUL-2004
05-JUL-2004
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Holden M.T.G., Feil E.J., Lindsay J.A., Peacock S.J., Day N.P.J.,
Enright M.C., Foster T.J., Moore C.E., Hurst L., Atkin R., Barron A.
Bason N., Bentley S.D., Chillingworth C., Chillingworth T.,
Churcher C., Clark L., Corton C., Cronin A., Doggett J., Dowd L.,
Feltwell T., Hance Z., Harris B., Hauser H., Holroyd S., Jagels K.,
James K.D., Lennard N., Line A., Mayes R., Moule S., Mungall K.,
Ormond D., Quail M.A., Rabbinowitsch E., Rutherford K.M., Sanders M.
Sharp S., Simmonds M., Stevens K., Whitehead S., Barrell B.G.,
Spratt B.G., Parkhill J.;
"Complete genomes of two clinical Staphylococcus aureus strains:
evidence for the rapid evolution of virulence and drug resistance.",
Proc. Natl. Acad. Sci. J.A. 101:9786-9791 (2004).
                                   COF2 HUMAN
Q9Y281;
30-MAY-2000
30-MAY-2000
25-OCT-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Staphylococcus aureus (strain MSSA476).
Bacteria; Firmicutes; Bacillales; Staphylococcus
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Putative surface anchored protein.
                                                                                                                                                                                                                                                                                                      EMBL; BX571857; CAG43485.1; -. GO; GO:000986; C:cell surface; IEA. GO; GO:0016020; C:membrane; IEA. InterPro; IPR011439; DUF1542.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OrderedLocusNames=SAS1682;
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Eukaryota; Metazoa;
          Homo sapiens (Human)
                    Name=CFL2;
                                                                                      HUMAN
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llarity 100.0%;
Conservative 0
                            (Rel. 39, Created)
(Rel. 39, Last sequence update)
(Rel. 45, Last annotation update)
scle isoform (Cofilin-2).
                                                                                                                                                                  Conservative
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                                                                            STANDARD;
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Chordata;
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Pred. No. 2.4e+03;
                                                                                                                                                                 Score 32; DB 2; ]
Pred. No. 2.4e+03;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRT;
 Craniata; Vertebrata;
                                                                             PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  621D7875D1D3FB35 CRC64;
                                                                                                                                                                                                         3A041AF93D45932C CRC64;
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                                                                                                                                                                                   Length 2186;
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  Euteleostomi;
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RA Strausberg R.D., Fedingold E.A., Grouse L.H., Derge J.G.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Schapteron M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Robak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Whiting M., Madan A., Rodrigues S., Sanchez A.,
RA Rodriguez A.C., Grimwood J., Schwutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
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RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimood R.M.,
RA Rodriguez A.C., Grimoo
                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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EMBL; AF134802; AAD31280.1;
EMBL; AF134803; AAD31281.1;
EMBL; AF283513; AAF97934.1;
EMBL; AF242299; AAF64498.1;
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"Isolation of two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=9606;
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                                                                                                                                                                           or send an
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Name=CFLZA;
IsoId=Q9Y281-2; Sequence=Not described;
TISSUE SPECIFICITY: Isoform CFL2b is expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity, AITERNATIVE PRODUCTS:
Event=Alternative splicing; Named isoforms=2;
Event=Alternative splicing; Named isoforms=2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SIMILARITY: Belongs to the actin-binding proteins ADF family.
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                                                                                                                                                                               email to license@isb-sib.ch).
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PubMed=12477932; DOI=10.1073/pnas.242603899;
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Peters G.J., Abramson R.D., Mullahy S.J.,
Srnan K.J., Malek J.A., Gunaratne P.H.,
le S., Garcia A.M., Gay L.J., Hulyk S.W.,
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EMBL; EMBL; EMBL; HSSP;

BC011444; BC022364; BC022876; P18282;

AAH11444.1; AAH22364.1; AAH22876.1;

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                                                                       Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan K., Farmer A.A., Rubin G.M., Hong L.,

A Hopkins R.F., Jordan R., Fonaldo M.A., Rubin G.M., Hong L.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Hillalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Pahey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Holkesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

A Hutterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

Ceneration and initial analysis of more than 15,000 full-length human
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Best Local S
Matches 5
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01-NOV-1995
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P45591;
and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002)
-!- FUNCTION: Controls reversibly actin polymeriza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DOMAIN
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SMART; SMOO102; ADF; 1.
PROSITE; P800325; ACTIN DEPOLYMERIZING; 1.
PROSITE; P800325; ATTACTIVE SPLICING; Cytoskeleton;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00241; Cofilin ADF; PRINTS; PR00006; COFILIN. ProDom; PD002129; Actbind_co
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MIM; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ono S. Minami N., Abe H., Obinata T.;
"Characterization of a novel cofilin isoform that is
expressed in mammalian skeletal muscle.";
J. Biol. Chem. 269:15280-15286(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=C3H; TISSUE=Skeletal musc1
MEDLINE=94253093; PubMed=8195165;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
STRAIN=C3H; TISSUE=Skeletal
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IGDIGD
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(Rel. 32, Last sequence up
(Rel. 45, Last annotation
uscle isoform (Cofilin-2).
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48B6CDCCAE9FE1CC;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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No. 2.7e+02
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Best Local &
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28-FEB-2003
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SEQUENCE
                                                                                                                                                                                  SEQUENCE FROM N.A.

TISSUB-Skeletal muscle;

MEDLINE=91027755, PubMed=1699599;

Abe H., Endo T., Yamamoto K., Obinata T.;

"Sequence of cDNAs encoding actin depolymerizing factor and cofilin embryonic chicken skeletal muscle: two functionally distinct actin-regulatory proteins exhibit high structural homology.";

Biochemistry 29:7420-7425(1990).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by an
                                                                                                               STRUCTURE BY NMR, AND REVISIONS TO MEDLINE=21880669; PubMed=11885570;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gallus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Actin-binding; Cytoskeleton; Nuclear protein; Phosphorylation.

DOMAIN

30

34

Nuclear localization signal (Potential).

DOMAIN

106

125

Actin-binding (Potential).

MOD RES

24

Phosphoserine (By similarity).

SEQUENCE

166

AA; 18709 MW; 48B6D7E5AE9FEICC CRC64;
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SMART; SM00102; ADF; 1.
PROSITE; PS00325; ACTIN DEPOLYMERIZING;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR002108; Actbind Pfam; PF00241; Cofilin ADF; PRINTS; PR00006; COFILIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; L29468; AAA37433.1; -.
EMBL; BC007138; AAH07138.1; -.
PIR; A53812; A53812.
                        cofilin.";
                                                                   Bains N.P.S., Gorl
Maciejewski M.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                 Gallus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Archosauria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota;
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                                               "Backbone and side-chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MGD; MGI:101763; Cfl2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         send
Biomol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       major component of intranuclear and cytoplasmic actin rods. SUBCELUTIAR LOCATION: Intranuclear and cytoplasmic TISSUE SPECIFICITY: Predominantly expressed in skeletal muscle. PTM: The phosphorylation of Ser-24 may prevent recognition of the nuclear localization signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SIMILARITY: Belongs to the actin-binding proteins ADF family.
                                                                                                                                                                                                                                                                                                                                                                                                   _TaxID=9031;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CHICK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gallus (Chicken).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VGDIGD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                requires a license agreement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Metazoa;
  NMR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (Rel. 18, Created)
(Rel. 41, Last seq
(Rel. 44, Last ann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            erization in a pH-sensitive manner. It and F-actin in a 1:1 ratio of cofilin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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Aves; Neognathae;
                                            NMR, AND REVISIONS TO 53-54.

669; PubMed=11885570; DOI=10.1023/A:1014227808686;

Gorbatyuk V.Y., Nosworthy N.J., Robson S.A.,

W., dos Remedios C.G., King G.F.;

side-chain 1H, 15N, and 13C assignments for chick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1AK6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Craniata; Vertebrata; Euteleostomi; 
; Galliformes; Phasianidae; Phasiani
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2.7e+02;
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assignments for chick
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Best Local S
Matches 5
                 Strausberg R.L., Feingold B.A., Grouse L.H., Derge J.G.,
Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
Brownstein M.J., McKernan K.J., Maramson R.D., Mullahy S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
The College C., College C., College C., College C., College C., Walle S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
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SEQUENCE
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MOD_RES
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                                                                                                                                                                                                                                                                                                                                                                                                                Brachydanio rerio (Zebrafish)
Eukaryota; Metazoa; Chordata;
Actinopterygii; Neopterygii; T
Cyprinidae; Danio.
                                                                                                                                                                                                                                                                                               TISSUE=Embryo;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Name=cfl2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
Cofilin 2 (Muscle).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Q6NZW3;
05-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Q6NZW3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRINTS; PR00006; COFILIN.
ProDom; PD002129; Actbind
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pfam; PF00241; Cofilin_ADF;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR002108; Actbind_cofln.
Pfam; PF00241; Cofilin_ADF; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nuclear localization signal.
SIMILARITY: Belongs to the actin-binding proteins ADF family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FUNCTION: Controls reversibly actin polymerization and depolymerization in a pH-sensitive manner. It has the ability bind G- and F-actin in a 1:1 ratio of cofilin to actin. It is major component of intranuclear and cytoplasmic actin rods:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PTM: The phosphorylation of Ser-24 may
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          dimethyl sulfoxide.
TISSUE SPECIFICITY: Widely_distributed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SUBCELLULAR LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             completely in nucleus in cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SM00102; ADF; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   μ
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5; Conserv
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     Helton E.,
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106
24
166 J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cytoskeleton;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACTIN_DEPOLYMERIZING; 1.
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125
24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Actbind_cofln;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18661 MW;
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83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .3%
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27,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Created)
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nuclear protein; Phosphorylation.
Nuclear localization signal (Potential).
Actin-binding (Potential).
Phosphoserine (By similarity).
; 514685E940786EF0 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               is produced through a c
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ng as its content is in
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2.7e+02;
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to heat shock or 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 166;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and the
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RESULT 18 Q9US43 ID Q9US4 AC Q9US4

Q9US43 Q9US43;

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RESULT 17
Q853Z5
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GO; GO:0005622; C:intracellular; II
GO; GO:000379; F:actin binding; II
InterPro; IPR002108; Actbind_cofln.
Pfam; PF00241; Cofilin_ADF; I.
PRINTS; PR00006; COFILIN.
ProDom; PD002129; Actbind_cofln; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUN-2003
01-JUN-2003
01-JUN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q853Z5
Q853Z5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ProDom; PD002129; Ac
SMART; SM00102; ADF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; BC065947; AAH65947.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Embryo;
Director MGC Project;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and mouse cDNA sequences."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Generation and initial analysis of more than 15,000 full-length human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jones S.J., Marra M.A.;
                                                                                                                                                                   "Origins of highly mosaic mycobacteriophage cell 113:171-182(2003).
EMBL, AY129338, AAN12813.1; -.
SEQUENCE 180 AA; 20106 MW; C7396B1AA20F0
                                                                                                                                                                                                                                                     Kriakov J., Lawrence J.G.,
Hatfull G.F.;
                                                                                                                                                                                                                                                                                                                   MEDIINE=22592660; PubMed=12705866; DOI=10.1016/S0092-8674(03)00233-2; Pedulla M.L., Ford M.E., Houtz J.M., Karthikeyan T., Wadsworth C., Lewis J.A., Jacobs-Sera D., Falbo J., Gross J., Pannunzio N.R.,
                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=205879;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Viruses;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mycobacteriophage Omega.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HSSP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (FEB-2004)
                                                                                                                                                                                                                                                                              Brucker W., Kumar V., Kandasamy J.,
Kriakov J., Lawrence J.G., Jacobs W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Name=171;
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PS00325; ACTIN DEPOLYMERIZING; 1.
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Smailus D.E., Schnerch A.,
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RA Sgouros J., Peat N., Hayles J., Baker S., Basham D., Bowman S.,
RA Brooks K., Brown D., Brown S., Chillingworth T., Churcher C.,
RA Brooks K., Brown D., Brown S., Chillingworth T., Churcher C.,
RA Collins M., Connor R., Cronin A., Davis P., Feltwell T., Fraser A.,
RA Collins M., Connor R., Cronin A., Davis P., Feltwell T., Fraser A.,
RA Holroyd S., Hornsby T., Howarth S., Huckle E.J., Hunt S., Jagels K.,
James K., Jones M., Leather S., McDonald S., McLean J.,
RA Holroyd S., Hornsby T., Howarth S., Huckle E.J., Rhut S., McLean J.,
RA Goliver K., O'Neil S., Pearson D., Quail M., Rabbinowitsch E.,
RA Oliver K., O'Neil S., Pearson D., Quail M., Rabbinowitsch E.,
RA Gulbert G., Rutter S., Saunders D., Seeger K., Sharp S.,
RA Taylor K., Taylor R.G., Tivey A., Walsh S., Warren T., Whitehead S.,
RA Moodward J., Volckaert G., Aert R., Robben J., Grymonprez B.,
RA Weltjens I., Vanstreels E., Rieger M., Schafer M., Muller-Auer S.,
RA Meltjens I., Vanstreels E., Rieger M., Schafer H., Reinhardt R.,
RA Hilbert H., Borzym K., Langer I., Beck A., Lehrach H., Reinhardt R.,
RA Hilbert H., Aves S.J., Xiang Z., Hunt C., Moore K., Hurst S.M.,
RA Galibert F., Aves S.J., Xiang Z., Hunt C., Moore K., Hurst S.M.,
RA Galibert F., Aves S.J., Xiang Z., Hunt C., Moore K., Hurst S.M.,
RA Dominguez A., Revuelta J.L., Moreno S., Armstrong J., Forsburg S.L.,
RA Cerutti L., Lowe T., McCombie W.R., Paulsen I., Potashkin J.,
RA Cerutti L., Lowe T., McCombie W.R., Paulsen I., Potashkin J.,
RT "The genome sequence of Schizosaccharomyces pombe.";
RT Nature 415:871-880(2002).
RE MBL; AL136078; CABS5617.1; -.
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Best Local (
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Q84BH1;
Q1-JUN-2003
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01-MAY-2000
01-MAR-2004
                                                                              01-JUN-2003 (TrEMBLrel. 24,
01-OCT-2003 (TrEMBLrel. 25,
Vls recombination cassette
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GeneDB SPombe; SPAC1002.17c; -.
GO; GO:001675; F.transferase activity, transferring glycosyl.
GO; GO:009316; p:nucleoside metabolism; IEA.
InterPro; IPR000508; Peptidase_S26.
InterPro; IPR00036; Peptidase_S26.
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01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Uracil phosphoribosyltransferase activity (Predicted); similar to
                             Borrelia afzelii.
Plasmid lp28-1.
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Pfam; PF00156; Pribosyltran; 1.
PROSITE; PS00501; SPASE_I_1; UNKNOWN_1.
Glycosyltransferase; Transferase.
Glycosyltransferase; Transferase.
SEOUENCE 189 AA; 20476 MW; A3930220404EC933 CRC64;
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Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
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Best Local Similarity
Matches 5; Conserv
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01-MAY-1999
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MEDLINE=22493136; PubMed=12603744;
Wang D., Botkin D.J., Norris S.J.;
"Characterization of the vls antigenic variation loci of the Lyme
"Characterization of arrelia gazinii Ip90 and Borrelia afzelii
                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=97275907; FUNDWEGE=122000,
Hu C.-C., Aboul-Ata A.E., Naidu R.A., Ghabrial S.A.;
"Evidence for the occurrence of two distinct subgroups of peanut
"Evidence for the occurrence of two distinct of RNA3.";
019062;
01-JAN-1998
                                                                                                                                                                                                                                                                  Pfam; PF00760; Cucumo coat; 1.
PRINTS; PR00222; CUCUMOCOAT.
ProDom; PD001284; Cucumovirus_coat; 1.
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GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule activity; IEA
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MEDLINE=97275907; PubMed=9129668;
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EMBL; AY100628; AAN87821.1; -.
HSSP; O68364; ILBW.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=29518;
[1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=12313;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cucumovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Viruses; ssRNA positive-strand viruses, no DNA stage; Bromoviridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peanut stunt virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Coat protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00921; Lipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                     InterPro; IPR000247; Cucumovirus_coat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GO:0016020; C:membrane; IEA.
                                                                                                                                                                                                                                                          protein
                                                                                                                               1 IGDIGD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 IGDIGD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                  Virol.
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                                                                                                           VGDIGD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IPR000680; Borrelia_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    201
201 AA;
                                                                                                                                                                                                                                         214 AA;
(TrEMBLrel. 05,
                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (TrEMBLrel. (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (TrEMBLrel.
                               PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                      78:929-939(1997).
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19522 MW;
                                                                                                                                                                                                                                         23765 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96.9%;
                                                                                                                                                                                         96.9%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 31; DB 2;
Pred. No. 3.3e+02;
1; Mismatches 0
Created)
                                                                                                                                                                          1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Created)
                                                                                                                                                                                           Score 31;
Pred. No.
                               PRT;
                                                                                                                                                                                                                                         9103CBB8BC0C9A93 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     174E3DF343708B01 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             214 AA.
                               222
                                                                                                                                                                                           3.6e+02;
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                                                                                                                                                                                                          Length 214;
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                                                                                                                                                                            Indels
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RESULT 22
CRP_CAVPO
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Rubio N., Sharp P.M., Rits M., Zahedi K., Whitehead A.&

"Structure, expression, and evolution of guinea pig ser

component and C-reactive protein.";

J. Biochem. 113:277-284(1993).

1-- FUNCTION. Displays several functions associated wit

it promotes agglutination, bacterial capsular swell

phagocytosi and complement fixation through its cal

binding to phosphorylcholine. Can interact with DNA

and may scavenge nuclear material released from dam

circulating cells (By similarity).

-!- SUBUNIT: Homopentamer. Pentaxin (or pentraxin) have

arrangement of 5 noncovalently bound subunits (By s:

-!- SUBCELLULAR LOCATION: Secreted.

-!- TISSUE SPECIFICITY: Found in plasma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 petection of SNPs and linkage and radiation hybrid map porcine C-reactive protein gene, CRP.; Submitted (AUG-2004) to the EMBL/GenBank/DDBJ databases. EMBL; AB005545; BAA21473.1; -. EMBL; AY014055; AAU13779.1; -. HSSP; P02741; 1LJ7.
InterPro; IPPn^^^
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CRP_CAVPO
P49254;
01-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Signal.
SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C-reactive protein precursor.
Sus scrofa (Pig).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CHAIN
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=Landrace;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JAN-1998 (TrEMBLrel. 05, Last 25-OCT-2004 (TrEMBLrel. 28, Last
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cavia porcellus (Guinea pig).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Hystricognathi; Caviidae; Cavia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ozawa A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C-reactive protein precursor.
Name=CRP; Synonyms=PTX1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-FEB-1996 (Rel. 33, Created)
01-FEB-1996 (Rel. 33, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SMART; SM00159; PTX;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRINTS; PR00895; PENTAXIN.
ProDom; PD002153; Pentaxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR000985; ConA_like_lec_gl.
InterPro; IPR001759; Pentaxin.
Pfam; PF00354; Pentaxin; 1.
                                                                                                                                                                                                                                                                                                                                      MEDLINE=93252746; PubMed=8486600;
                                                                                                                                                                                                                                                                                                                                                                 STRAIN=Hartley;
                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=10141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 169
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5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ignign 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VGDIGD 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PS00289;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matsumoto M., Kajikawa (JUL-1997) to the EMBL/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1
20
222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PENTAXIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24924 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ikawa M., Hanazono
EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ۲.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Potential.
CRP mature peptide.
; 451B7959DD5F8E9B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence update)
annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B
                                                                                                                                                                                                                       associated with host defense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .7e+02;
                                                                                                                                                                                                    capsular swelling,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      M., Yasue
databases.
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                                                                                   have a discoid
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Sus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mapping
                                                                                                                                                                                                                                                                                            A.S.;
serum
                                                                                                                                                   calcium-dependent
DNA and histones
                                                                                                                                 damaged
                                                             similarity)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            H.;
                                                                                                                                                                                                                                                                                              amyloid
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RESULT 23
CRP_MESAU
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     Query Match
Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           DISULFID
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-sor send an email to license@isb-sib.ch).
                                                                                                                                                                                                           NCBI_TaxID=10036;
                                                                                                                                                                                                                                                     Name=CRP; Synonyms=PTX1;
Mesocricetus auratus (Go
                                                                                                                                                                                                                                                                         C-reactive protein precursor.
                                                                                                                                                                                                                                                                                 05-JUL-2004 (Rel.
                                                                                                                                                                                                                                                                                            01-FEB-1996
01-FEB-1996
                                                                                                                                                                                                                                                                                                                 P49262;
                                                                                                                                                                                                                                                                                                                            CRP_MESAU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SMART; SM00159; PTX; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PIR; JX0259; JX0259.
HSSP; P02741; 1B09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; S60422; AAC60662.1; -.
                                                                                                                                                            Dowton S.B., Holden S.N.; "C-reactive protein (CRP) of the Syrian hamster.";
                                                                                                                                                                                MEDLINE=91369977; PubMed=1892852;
                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                        Mesocricetus
                                                                                                                                                                                                                                 Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Acute phase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ProDom; PD002153; Pentaxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRINTS; PR00895; PENTAXIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00354; Pentaxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR001759; Pentaxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR008985;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SIMILARITY: Belongs to the pentaxin family.
DATABASE: NAME=Protein Spotlight; NOTE=Issue 30 of January 2003;
WWW-"http://www.expasy.org/spotlight/articles/sptlt030.html".
                                                                                                                                                                                                                                                                                                                                                                           172 VGDIGD 177
                                                                                                                                                                                                                                                                                                                                                                                                1 IGDIGD 6
                                                                                                                                                                                                                                                                                                                                                                                                                     <u>ن</u>
                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PS00289; PENTAXIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             225
                                                                                                                                                                                                                                                                                            (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Calcium;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ΑA;
                                                                                                                                                                                                                                                                                                                            STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   19
225
225
116
116
25225 N
                                                                                                                                                                                                                                                                                             33,
33,
                                                                                                                                                                                                                                                                                    44, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                             96.9%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ConA_like_lec_gl.
                                                                                                                                                                                                                                                     (Golden hamster)
                                                                                                                                                                                                                                                                                                         Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pentaxin;
                                                                                                                                                                                                                                                                                              Last
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                                                                                                                                                                                                                                                                                             sequence
                                                                                                                                                                                                                                                                                                                                                                                                                     1;
                                                                                                                                                                                                                                                                                                                                                                                                                    Score 31; DB 1;
Pred. No. 3.7e+02;
1; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      By similarity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               By similarity.
C-reactive protein
Pentaxin.
                                                                                                                                                                                                                                             Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             47AF3E2D9A660F2E CRC64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Plasma; Signal.
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                                                                                                                                                                                                                                                                                                                             225 AA
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                                                                                                                                                                                                                                                                                    update)
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RESULT 24
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                                                                                                                                                              SEQUENCE FROM N.A.

STRAIN=CBA/J; TISSUE=Liver;

MEDLINE=90179719; pubMed=2310378;

Whitehead A.S., Zahedi K., Rits M., Mortensen R.F., Lelias J.M.;

"Mouse C-reactive protein. Generation of cDNA clones, structural analysis, and induction of mRNA during inflammation.";

Biochem. J. 266:283-290(1990).

-I- FUNCTION: Displays several functions associated with host defense it promotes agglutination, bacterial capsular swelling, phagocytosi and complement fixation through its calcium-dependent binding to phosphorylcholine. Can interact with DNA and histones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CRP_MOUSE
P14847;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DISULFID SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-APR-1990 (Rel. 14, Last sequence update) 05-JUL-2004 (Rel. 44, Last annotation updat C-reactive protein precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRINTS; PR00895; PENTAXIN.
ProDom; PD002153; Pentaxin; 1.
SMART; SM00159; PTX; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; S56005; AAB19893.2;
PIR; A40326; A40326.
HSSP; P02741; 1LJ7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=89050112; PubMed=3190681; Ohnishi S., Maeda S., Nishiguchi S., Ohnishi S., Maeda S., Nishiguchi S., Ohnishi S., Distructure of the mouse C-reactive pi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR008985; ConA like InterPro; IPR001759; Pentaxin. Pfam; PF00354; Pentaxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Name=Crp; Synonyms=Ptx1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                or send an email to license@isb-sib.ch).
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                       and may scavenge nuclear material rel
circulating cells.
SUBUNIT: Homopentamer. Pentaxin (or g
arrangement of 5 noncovalently bound
SUBCELLULAR LOCATION: Secreted.
TISSUB SPECIFICITY: Found in plasma.
INDUCTION: The concentration of CRP i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 IGDIGD
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5; Conserv
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225
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24822
                       CITY: Found in plasma. concentration of CRP in
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By similarity.
; 00B439C45D58E05A CRC64;
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Pred. No.
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C-reactive protein
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tive protein gene.";
156:814-822(1988).
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3.7e+02;
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n plasma
injury,
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    increases infection
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or other
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RESULT 25
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RX MEDILINE-2238855; Pubmedeil2477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold B.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altechul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
WA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
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Best Local
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CONFLICT
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                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                   Name=Crp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Q91XB3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q91XB3
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InterPro; IPR008985; ConA like
InterPro; IPR001759; Pentaxin.
Pfam; PF00354; Pentaxin; 1.
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                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
STRAIN=FVB/N; TISSUE=Liver;
                                                                                                                                                                                                                                                                                     NCBI_TaxID=10090;
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MGD; MGI:88512; Crp
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SIMILARITY: Belongs to the pentaxin family.
DATABASE: NAME=Protein Spotlight; NOTE=Issue 30 of January 2003;
WWW-"http://www.expasy.org/spotlight/articles/sptlt030.html".
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              172
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Rodentia;
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Best Local :
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MGD; MGI:88512; Crp.
GO; GO:0005615; C:extracellular space; TA
Pfam; PF00354; Pentaxin; 1.
PRINTS; PR00895; PENTAXIN.
ProDom; PD002153; Pentaxin; 1.
SMART; SM00159; PTX; 1.
SMART; SM00159; PTX; 1.
PROSITE; PS00289; PENTAXIN; 1.
SEQUENCE 225 AA; 25387 MW; 0E5FD795E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CRP_RAT
P48199;
                                                             it promotes agglutination, bacterial capsular swelling, phagocytosi and complement fixation through its calcium-dependent binding to phosphorylcholine. Can interact with DNA and histones and may scavenge nuclear material released from damaged circulating cells (By similarity).

-I SUBUNIT: Homopentamer. Pentaxin (or pentraxin) have a discoid arrangement of 5 noncovalently bound submits. Two of the five chains form a dimer linked by two interchain disulfide bonds located in the C-terminal heptapeptide and specific to rat CRP.
-I SUBCELULAR LOCATION: Secreted.
-I TISSUE SPECIFICITY: Found in plasma.
-I PTM: The last two cysteines are involved either in interchain disulfide bonds or in an intrachain bond of January 2003; WMW="http://www.expasy.org/spotlight/NOTE=Issue 30 of January 2003; WMW="http://www.expasy.org/spotlight/articles/sptlt030.html".
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nagpurkar A., Mookerjea S., Murray R.K.;
"Derivation of the amino acid sequence of rat C-reactive protein in the nature of its dimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rassouli M., Sambasivam H., Azadi P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=Sprague-Dawley; TISSUE-Liver; MEDLINE=92147634; PubMed=1737750;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rattus norvegicus (Rat).
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05-JUL-2004 (Rel. 44, Last annotation update)
C-reactive protein precursor.
Name=Crp; Synonyms=Ptx1;
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STRAIN=FVB/N; TISSUE=Liver;
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Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Component
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l. Acad. Sci. U.S
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Sciurognathi; Muridae; Murinae; Rattus.
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the European Bioinformatics Institute. There are no restrictions on its
use by non-profit institutions as long as its content is in no way
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entitles requires a license agreement (See http://www.isb-sib.ch/announce/
or send an email to license@isb-sib.ch).
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SMART; SM00159; PTX; 1.
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PRINTS; PR00895; PENTAXIN.
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GlycoSuiteDB; P48199;
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                                                                                                                                                                                                                          MEDILINE=22435905; PubMed=12547188; DOI=10.1016/S0022-2836(02)01400-6; Maeda K., Nojiri H., Shintani M., Yoshida T., Habe H., Omori T.; Complete nucleotide sequence of carbazole/dioxin-degrading plasmid pCAR1 in Pseudomonas resinovorans strain CA10 indicates its mosaicity and the presence of large catabolic transposon Tn4676."; J. Mol. Biol. 326:21-33(2003).
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Acute phase; Calcium; Direct
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InterPro; IPR001759; Pentaxin.
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GO; GO:0008152; P:metabolism; IEA.
InterPro; IPR002198; ADH_short.
InterPro; IPR002347; Adh_short_C2.
Pfam; PF00106; adh_short; 1.
PRINTS; PR000081; GDHRDH.
PRINTS; PR000081; GDHRDH.
PRINTS; PR00080; SDRFAMILY.
PROSITE; PS00061; ADH_SHORT; 1.
Oxidoreductase.
Hardenack S., De Y., Koutnikova F. Submitted (APR-1996) to the EMBL/-1- SIMILARITY: Belongs to the sh (SDR) family.

EMBL; US3828; AAB42054.1; -.
HSSP; P25716; 1Q7B.
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01-NOV-1996
01-NOV-1996
01-MAR-2004
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-!- SIMILARITY: Belongs to the short-chain dehydrogenases/reductases
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                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                      NCBI_TaxID=52853;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Name=STA1-12;
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Pred. No. 4.2e+02;
1; Mismatches 0
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CCD5 CAEEL

' P91285;

f 01-NOV-1997 (Rel. 35, Created)

T 01-NOV-1997 (Rel. 35, Last sequence updat

T 25-OCT-2004 (Rel. 45, Last annotation upr

DE Cuticle collagen dpy-5 (Dumpy protein 5)

Name=dpy-5; ORFNames=F27C1.8;
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Best Local S
Matches 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GO; GO:0016491; F:oxidoreductase act
GO; GO:0008152; P:metabolism; IEA.
InterPro; IPR002198; ADH short.
InterPro; IPR002347; Adh_short_C2.
Pfam; PF00106; adh short; 1.
PRINTS; PR00081; GDHRDH.
PRINTS; PR00080; SDRFAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
STA1-18 (STA1-2).
Name=STA1-18; Synonyms=STA1-2;
Silene pratensis (White Campion) (Lychnis alba).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Caryophyllales; Caryophyllaceae; Silene.
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InterPro; IPR002198; ADH_short.
InterPro; IPR002347; Adh_short_C2.
Pfam; PF00106; adh short; 1.
PRINTS; PR00081; GDHRDH.
PRINTS; PR00080; SDRFAMILY.
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EMBL; U53827; AAB42053.1; -.
HSSP; P25716; 107B.
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Hardenack S., De Y., Kou
Submitted (APR-1996) to
-!- SIMILARITY: Belongs
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01-NOV-1996
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SEQUENCE 281 AA;
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5; Conserve
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83.3%;
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83.3%;
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to the EMBL/GenBank/DDBJ databases.
1gs to the short-chain dehydrogenase
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Last sequence up
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                      Score 31; DB 2;
Pred. No. 4.7e+02
1; Mismatches
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Best Local
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Q8DGV6;
Q8DGV6;
Q1-MAR-2003 (TIEMBLICE). 2
Q1-MAR-2003 (TIEMBLICE). 2
Q1-OCT-2003 (TIEMBLICE). 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
STRAIN=Bristol N2;
MEDLINE=99069613; PubMed=9851916;
                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR008161; Clg_helix.
InterPro; IPR008160; Col_cuticle_N.
InterPro; IPR008160; ColTagen.
InterPro; IPR008160; ColTagen.
Pfam; PF01484; Col cuticle_N; 1.
Pfam; PF01391; ColTagen; 2.
ProDom; PD000007; Clg_helix; 2.
ProDom; PD000007; Clg_helix; 2.
Collagen; Cuticle; Multigene family; Repeat; Structural Collagen; Cuticle; Multigene family; Repeat; Structural DOMAIN
126
Triple-helical region.
DOMAIN
143
270
Triple-helical region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=6239;
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Rhabditidae; Peloderinae; Caenorhabditis.
                                                                                Synechococcus elongatus (Thermosynechococcus elongatus). Bacteria; Cyanobacteria; Chroococcales; Synechococcus.
                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "dpy-5 encodes a cuticle collagen.";

(er) Worm Breeder's Gazette 15(1):54(1997).

-I- FUNCTION: Nematode cuticles are composed largely of collagen-like proteins. The cuticle functions both as an exoskeleton and as a barrier to protect the worm from its environment.

-I- SUBUNIT: Collagen polypeptide chains are complexed within the cuticle by disulfide bonds and other types of covalent cross-links.
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Science 282:2012-2018(1998).
                                                                                                                 OrderedLocusNames=tlr2206;
                                                                                                                                UDP-glucose 4-epimerase
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WormPep; F27C1.8; CE09720.
 MEDLINE=22225144; PubMed=12240834;
                  SEQUENCE FROM N.A.
STRAIN=BP-1;
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                                                             NCBI_TaxID=32046;
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Pred. No.
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RESULT 34
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GO; GO:0003978; F:UDP-glucose 4-epimerase activity; IEA.
GO; GO:0006012; P:galactose metabolism; IEA.
GO; GO:0009225; P:nucleotide-sugar metabolism; IEA.
InterPro; IPR001509; Epimerase_Dh.
InterPro; IPR001509; Epimerase_Dh.
Ffam; PF01370; Epimerase; 1.
TIGRFAMB; TIGR01179; galE; 1.
COMPLETE Proteome; NAD.
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Morimura K., Ikeda H., Hat
"Complete genome sequence
thermophilum.";
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Bacteria; Actinobacteria; Sym
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1- SIMILARITY: Belongs to the sugar epimerase family.
EMBL, AP005376; BAC09758.1; -.
HSSP; Q8T8E9; 1GY8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chermophitum.";
Submitted (APR-2004) to the EMBL/GenBank/DDBJ databases.
... crwff.APTTV: Relongs to the sugar epimerase family.
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InterPro; IPR005886; GalE.
Pfam; PP01370; Epimerase; 1.
TIGRFAMs; TIGR01179; galE; 1.
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Hattori M., Beppu T.;
nce of an uncultured bacterium Symbiobacterium
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1; Mismatches
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Pred. No. 5.5e+02;
1; Mismatches 0
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Thomson N.R., Chatter K.F., Cerdeno-Tarraga A.-M., Kieser H.,

Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth S.,

Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth S.,

Huang C.-H., Kieser T., Larke L., Murphy L.D., Oliver K., O'Neil S.,

Rabbinowitsch E., Rajandream M.A., Rutherford K.M., Rutter S.,

Seeger K., Saunders D., Sharp S., Squares R., Squares S., Taylor K.,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=21477403; PubMed=11572948; DOI=10.1073/pnas.211433198; Omura S., Kheda H., Ishikawa J., Hanamoto A., Takahashi C., Shinose M., Takahashi Y., Horikawa H., Nakazawa H., Osonoe T., Kikuchi H., Shiba T., Sakaki Y., Hattori M.; "Genome sequence of an industrial microorganism Streptomyces "Genome sequence of an industrial microorganism Streptomyces avermitilis: deducing the ability of producing secondary
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Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
                                       "Complete genome sequence of the model coelicolor A3(2).";
Nature 417:141-147 (2002).
EMBL; AL939119; CAC08313.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales; Streptomycineae; Streptomycetaceae; Streptomyces.
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01-JUN-2003
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tegral membrane protein.
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   Q7XTS7;
01-OCT-2003
01-MAR-2004
01-MAR-2004
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01-MAY-2000
01-MAY-2000
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MEDLINB=20036896; PubMed=10567266; DOI=10.1126/Science.286.5444.1571;
White O, Eisen J.A., Heidelberg J.F., Hickey E.K., Peterson J.D.,
Dodson R.J., Haft D.H., Gwinn M.L., Nelson W.C., Richardson D.L.,
Moffat K.S., Qin H., Jiang L., Pamphile W., Crosby M., Shen M.,
Vamathevan J.J., Lam P., McDonald L.A., Utterback T.R., Zalewski C.,
Makarova K.S., Aravind L., Daly M.J., Minton K.W., Pleischmann R.D.,
Ketchum K.A., Nelson K.E., Salzberg S.L., Smith H.O., Venter J.C.,
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01-MAY-2000 (TrEMBLrel. 13, Last sequence u
01-MAR-2004 (TrEMBLrel. 26, Last annotation
Thymidine diphosphoglucose 4,6-dehydratase.
OrderedLocusNames-DRA0041;
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                                                                                                                                          Q7XTS7
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InterPro; IPR005888; dTDF gluc dehyt.
InterPro; IPR001509; Epimerase Dh.
Pfam; PF01370; Epimerase; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Science 286:1571-1577(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Genome sequence of the radioresistant bacterium Deinococcus radiodurans R1.";
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5; Conserv
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P:nucleotide-sugar metabolism;
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Last annotation update)
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Pred. No. 5.9e+02;
1; Mismatches 0;
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SEQUENCE FROM N.A.
STRAIN=MAFF303099;
STRAIN=MAFF303099;
MEDLINE=21082930; PubMed=11214968;
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Matanabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura Kishida Y., Kiyokawa C., Kohara M., Matsumoto M., Matsuno Mochizuki Y., Nakayama S., Nakazaki N., Shimpo S., Sugimot Takeuchi C., Yamada M., Tabata S.;
Takeuchi C., Yamada M., Tabata S.;
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Liu Y., Hu X., Jia P., Zhang Y., Zhao Q., Ying K., Yu S., Tang Y.,
Weng Q., Zhang L., Lu Y., Mu J., Lu Y., Zhang L.S., Yu Z., Fan D.,
Liu X., Lu T., Li C., Wu Y., Sun T., Lei H., Li T., Hu H., Guan J.,
Wu M., Zhang R., Zhou B., Chen Z., Chen L., Jin Z., Wang R., Yin H.,
Cai Z., Ren S., Lv G., Gu W., Zhu G., Tu Y., Jia J., Zhang Y.,
Chen J., Xang H., Chen X., Shao C., Sun Y., Hu Q., Zhang X., Zhang W.
Wang L., Ding C., Sheng H., Gu J., Chen S., Ni L., Zhu F., Chen W.,
Lan L., Lai Y., Cheng Z., Gu M., Jiang J., Li J., Hong G., Xue Y.,
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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01-OCT-2001 (TrEMBLrel. 18, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation updat
NADH-ubiquinone dehydrogenase chain 3.
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Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales;
Phyllobacteriaceae; Mesorhizobium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OrderedLocusNames=mll1362;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR001878; Znf_CCHC.
SMART; SM00343; Znf_C2HC; 1.
SEQUENCE 588 AA; 64054 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Sequence and analysis of rice chromosome 4."; Nature 420:316-320(2002).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PubMed=12447439;
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cathity):

CATALTIC ACTIVITY: NADH + quinone = NAD(+) +
SIMILARITY: Belongs to the complex I 75 kDa su
L; AP002997; BAB48752.1; -
GO:0005489; F:electron transporter activity; I
GO:0005506; F:iron ion binding; IEA.
GO:0008137; F:NADH dehydrogenase (ubiquinone)
GO:0016651; F:oxidoreductase activity, acting
                                                                                                                                                                                                                                         Res. 7:331-338(2000). FUNCTION: NDH-1 shutt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GO:0003676; F:nucleic acid binding;
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        activity; on NADH or
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chain (By
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GYRB_RICCN
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Best Local
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Q92H87;
                                                                                                                                             collabor muticular institute. There are no restrictions on the statement is not removed. Usage by and for commerce are during a license agreement (See http://www.isb-sib.ch/~~send an email to license@isb-sib.ch).
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NAD; Oxidoreductase; Quinone; Ubiquinone.
PROTENCE 693 AA; 74544 MW; 6CC39A61B38A7631 CRC64;
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Pfam; PF00314; Molybdopterin; 1.
TIGREAMS; TIGR01973; NuoG; 1.
PROSITE; PS00642; COMPLEX1 75K 2; 1.
PROSITE; PS00643; COMPLEX1 75K 3; 1.
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InterPro; IPR001041; Ferre
InterPro; IPR006656; Molyb
InterPro; IPR010228; NuoG.
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28-FEB-2003 (Rel.
05-JUL-2004 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=21442074; PubMed=11557893; DOI=10.1126/science.1061471; Ogata H., Audic S., Renesto-Audiffren P., Fournier P.-E., Barbe Samson D., Roux V., Cossart P., Weissenbach J., Claverie J.-M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiaceae; Rickettsieae; Rickettsia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rickettsia conorii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA gyrase subunit B (EC 5.99.1.3).
Name=gyrB; OrderedLocusNames=RC0884;
                                                                                            HSSP; P06982; 1AJ6.
InterPro; IPR003594;
                                                                                                                                           EMBL; AE008642; AAL03422.1; PIR; D97810; D97810.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA rings, including catenanes and knotted rings. CATALYTIC ACTIVITY: ATP-dependent breakage, passa of double-stranded DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                  SIMILARITY: Belongs to the type II topoisomerase family.
                                                                                                                                                                                                                                                                                                                                                                                                                                         breakage and rejoining; the B chain enzyme forms an A2B2 tetramer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUBUNIT: Made up of two chains. The A chain is breakage and rejoining; the B chain catalyzes A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FUNCTION: DNA gyrase negatively supercoils closed circular double-stranded DNA in an ATP-dependent manner and also catalyzes the interconversion of other topological isomers of double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  _TaxID=781;
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0283; Complex1 75K.
1041; Ferredoxin.
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L B (EC 5.99.1.3).
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83.3%;
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ATP hydrolysis.
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nterPro;

IPR000565; IPR001241; IPR002288;

; ATPbind_ATPase.; DNA_gyraseB_C.; DNA_gyrB.; DNA_topoisoII.; Toprim_dom.

interPro; nterPro;

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Pfam; PF00986; DNA_gyraseBC; 1.
Pfam; PF02518; HATPase C; 1.
Pfam; PF01751; Toprim; 1.
PfNTS; PR00418; TP12FAMILY.
PRINTS; PR00418; TP12FAMILY.
PRODOm; PD149633; DNA_gyrase_B; 1
SMART; SM00387; HATPase_C; 1.
                                                                                                                                                                                                                                                        entities
or send a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE-99039499; PubMed-9823893; DOI=10.1038/24094; Andersson S.G.E., Zomorodipour A., Andersson J.O., Sicheritz-Ponten T., Alsmark U.C.M., Podowski R.M., Naeslund Eriksson A.-S., Winkler H.H., Kurland C.G.; "The genome sequence of Rickettsia prowazekii and the origin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q9ZCX2;
30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence up
05-JUL-2004 (Rel. 44, Last annotation
DNA gyrase subunit B (EC 5.99.1.3).
Name-gyrB; OrderedLocusNames=RP580;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TIGRRAMS; TIGRO1059; GYTB; 1.
PROSITE; PS00177; TOPOISOMERASE II; 1.
ATP-binding; Complete procheome; Isomerase; Topoisomerase.
SEQUENCE 807 AA; 90794 MW; 12814B49F12132F9 CRC64;
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SMART; SM00433; TOP2c; 1
                                                                                                                                                                                                                                                                                                           use by non-profit institutions as long a modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nature 396:133-140(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales; Rickettsiaceae; Rickettsieae; Rickettsia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GYRB RICPR
  InterPro; IPRC
Pfam; PF00204;
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                                                                               [nterPro;
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                                                    nterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          s SWISS-PROT entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation - Buropean Bioinformatics Institute. There are no restrictions on its by non-profit institutions as long as its content is in no way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FUNCTION: DNA gyrase negatively supercoils closed circular double-
stranded DNA in an ATP-dependent manner and also catalyzes the
interconversion of other topological isomers of double-stranded
DNA rings, including caten
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIMILARITY: Belongs to the type II topoisomerase family
                                                                                                                                                      ,; AJ235272; CAA1
A71663; A71663.
); P06982; 1AJ6.
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                       ; IPR003594; ; iPR002288; ; iPR000565; ; iPR0001241; ; iPR001241; ; iPR006171;
                                                                                                                                                                                                                                                        requires a license agreement (Some an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IGDVGD
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                                                                                                                                                                                                      CAA15027.1;
                                             ; ATPbind ATPase.
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; DNA_gyrB.
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Pred. No. 1.4e+03;
1; Mismatches (
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                       Query Match
  Best Local Similarity
                                                                                                                                         InterPro; IPR003594; ATPBIND ATPASE.
InterPro; IPR00228; DNA GYYTASEB C.
InterPro; IPR011558; DNA GYYTASEB C.
InterPro; IPR011557; GYTB.
InterPro; IPR011557; GYTB.
InterPro; IPR00171; Toprim dom.
Pfam; PF00204; DNA GYYTASEB C; 1.
Pfam; PF00206; DNA GYTASEB; 1.
Pfam; PF00206; DNA GYTASEB C; 1.
Pfam; PF01518; HATPASE C; 1.
Pfam; PF01518; HATPASE C; 1.
Pfam; PF01518; TOPPIM; 1.
PRINTS; PR00418; TPIZERMILY.
PTODOM; PD149633; DNA GYTASE B; 1.
TIGREAMS; TIGR01059; GYTB; 1.
TIGREAMS; TIGR01059; GYTB; 1.
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Pfam; PF02518; HATPase C; 1.
Pfam; PF01751; Toprim; 1.
PRINTS; PR00418; TP12FAMILY.
PRODOM; PD149633; DNA_gyrase_B; 1
SMART; SM00387; HATPase C; 1.
SMART; SM00433; TOP2C; 1.
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Q7P921;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Malek J.A., Eremeeva M.E., Dasch G.A.;
Submitted (FEB-2003) to the EMBL/GenBank/DDBJ databases.
Submitted (FEB-2003) to the EMBL/GenBank/DDBJ databases.
-i- FUNCTION: DNA gyrase negatively supercoils closed circular double stranded DNA in an ATP-dependent manner and also catalyzes the interconversion of other topological isomers of double-stranded DNA rings, including catenanes and knotted rings (By similarity) DNA rings.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TIGRFAMS; TIGR01059; gyrB; 1.
PROSITE; PS00177; TOPOISOMERASE_II; 1.
ATP-binding; Complete proteome; Isomer
SEQUENCE 807 AA; 90823 MW; 5DED5A8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HSSP;
GO; G
GO; G
GO; G
                                                                           ATP-binding; Isomerase; Topoisomerase SEQUENCE 807 AA; 90651 MW; 53FED1/
                                                                                                                    TIGRFAMS; TIGR01059; gyrB; 1.
PROSITE; PS00177; TOPOISOMERASE II;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rickettsiaceae; F
NCBI_TaxID=35793;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Name=rsib_orf.1216;
Rickettsia_sibirica.
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01-MAR-2004
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GO:0005524; F:ANP binding; IEA.
GO:0003918; F:DNA topoisomerase (ATP-hydro
GO:0006304; P:DNA modification; IEA.
GO:0006365; P:DNA topological change; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of double-stranded DNA.
SIMILARITY: Belongs to the type II topoisomerase family.
CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      preliminary data.
, AABW01000001; EAA26372.1;
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5; Conser
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Last annotation update)
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Pred. No.
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                                                                             53FED1A519AF358A CRC64;
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     1.4e+03;
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RESULT 43
Q7TMA9
ID Q7TMA
AC Q7TMA
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Y. Bacteriol. 186:5842-5855 (2004).

CC :- FUNCTION: DNA gyrase negatively supercoils closed circular double-
cc stranded DNA in an ATP-dependent manner and also catalyzes the
cc interconversion of other topological isomers of double-stranded
CC :- CATALYTIC ACTIVITY: ATP-dependent breakage, passage and rejoining
CC :- SIMILARITY: Belongs to the type II topoisomerase family.

CC :- SIMILARITY: Belongs to the type II topoisomerase family.

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ATP-binding;
SEQUENCE 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PubMed=15317790; DOI=10.1128/JB.186.17.5842-5855.2004; McLeod M.P., Qin X., Karpathy S.E., Gioia J., Highlander S.K., Fox G.E., McNeill T.Z., Jiang H., Muzny D., Jacob L.S., Hawes Sodergren E., Gill R., Hume J., Morgan M., Fan G., Amin A.G., Gibbs R.A., Hong C., Yu X.-J., Walker D.H., Weinstock G.M.; "Complete genome sequence of Rickettsia typhi and comparison we sequences of other Rickettsiae.";
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5.99.1
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25-OCT-2004
25-OCT-2004
  Q7TMA9
Q7TMA9;
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PubMed=15317790; D
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rickettsiaceae;
NCBI_TaxID=785;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bacteria; Proteobacteria; Alphaproteobacteria;
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                                                                                                                                                                                                                                                                                                                                                                  TIGRFAMs;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     42
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                                                                                                                                                                                                                Similarity 5; Conser
                                                                                                                                                                                                                                                                                             ; TIGRO1059; GYRB; 1.
PS00177; TOPOISOMERASE_II; 1.
ing; Complete proteome; Isomerase; Topoisomerase.
807 AA; 90951 MW; 659227788BF99FF2 CRC64;
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(TrEMBLrel. 28, I
merase II subunit
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                                                                                                                                                                                                                   Conservative
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28,
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t B (DNA gyrase subunit
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                                                                                       Query Match
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Matches 5
                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.

Xu C.S., Li W.Q., Li Y.C., Yang K.J., Yan H.M., (Mang L., Wang S.F., Han H.P., Wang G.P., (Ma H., Wang L., Wang S.F., Han H.P., Wang G.P., (Ma H., Wang L., Wang Q.N., Zhang J.B.;
Shi J.B., Rahman S., Wang Q.N., Zhang J.B.;
Shi J.B., Rahman S., Wang Q.N., Zhang J.B.;
Submitted (JUN-2003) to the EMBL/GenBank/DDBJ datement of the Company of the EMBL/GenBank/DDBJ datement of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of the Company of th
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XU C.S., Li W.Q., Li Y.

Yang K.J., Yan H.M., C.

Shi J.B., Rahman S., W.

Submitted (JUN-2003) t
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01-OCT-2003 (TrEM)
05-JUL-2004 (TrEM)
Ac1262 (Ac1-114)
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Xu C.S., Li W.Q., Li Y.C., Chang C.F., Han H.F.

Xu C.S., Zhi W.Q., Li Y.C., Chang C.F., Man Yuan J.Y., Yang K.J., Yan H.M., Zhao L.F., Ma Shi J.B., Rahman S., Wang Q.N., Zhang J.B.; Submitted (JUN-2003) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.

XU C.S., Li W.Q., Li Y.C., Wang L., Wang S.F., Han H.P.,

XU C.S., Yuan J.Y., Yang K.J., Yan H.M., Chang C.F., 2

Chai L.Q., Yuan J.Y., Yang K.J., Yan H.M., Chang C.F., 2

Ma H., Shi J.B., Rahman S., Wang Q.N., Zhang J.B.;

Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases
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Xu C.S., Li W.Q., Li Y.C., Han H.P., W
Xu G.S., Yan H.M., Chang C.F., Zhao
Yang K.J., Yan H.M., Chang Q.N., Zhang
Shi J.B., Rahman S., Wang Q.N., Zhang
Submitted (MAY-2003) to the EMBL/GenBa
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Mammalia; Eutheria;
NCBI_TaxID=10116;
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XU C.S., Li W.Q., Li Y.C., Zhao L.F., Ma H., Wang L., Wa Yu C.B., Wang G.P., Chai L.Q., Yuan J.Y., Yang K.J., Ya Han H.P., Wang G.P., Chai J.B., Rahman S., Wang Q.N., Zhang J.B.; Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.
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Chang C.F., Zhao L.F., Ma H., Wang L., Wang C.
Chai L.Q., Shi J.B., Rahman S., Wang Q.N., Zha
Submitted (JUN-2003) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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SMART; SM00159;
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  h 96.9%;
Similarity 83.3%;
5; Conservative
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                                                                                         PS00213; LIPOCALIN;
PS00289; PENTAXIN;
1015 AA; 116461 I
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1 (Ac2-069) (J
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Da; Chordata;
ia; Rodentia;
                                                                                                                                                                            Pentaxin;
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C.F., Zhao L.F., Ma
Q.N., Zhang J.B.;
e EMBL/GenBank/DDBJ d
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(Ab2-196) (Ab1-341) (Ba2
      Score
Pred.
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Sciurognathi; Muridae;
           Mismatches
                                                                                              94904AC6BB979A76 CRC64;
                               31;
No.
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databases

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Wang

(Ba2-693)

Euteleostomi; ; Murinae; Rati

Rattus

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Wang

.8e+03; 0; DB 2;

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Gaps

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Length 1015;

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788 VGDIGD

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STRAIN=C57BL/6J; TISSUE=Heart;

MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100;

Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K.,

Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;

"Normalization and subtraction of cap-trapper-selected cDNAs

prepare full-length cDNA libraries for rapid discovery of new

Genome Res. 10:1617-1630(2000).
STRAIN=C57BL/6J; TISSUE=Heart;
STRAIN=C57BL/6J; TISSUE=Heart;
Adachi J., Aizawa K., Akimura T., Arakawa T., Bono H., Ca:
Fukuda S., Furuno M., Hanagaki T., Hara A., Hashizume W.,
Fukuda S., Furuno M., Hiramoto K., Hiraoka T., Hiroza
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01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Mus musculus 13 days embryo heart cDNA, RIKEN full-length enriched
library, clone:D330046B13 product:minichromosomo maintain
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STRAIN=C57BL/6J; TISSUE=Heart;
MEDLLNE=21085660; PubMed=11217851; DOI=10.1038/35055500;
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MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
Carninci P., Hayashizaki Y.;
"High-efficiency full-length cDNA cloning.";
Meth. Enzymol. 303:19-44(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the RIKEN Genome Exploration Research Group Phase I & II Team; "Analysis of the mouse transcriptome based on functional annotation 60,770 full-length cDNAs.";
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Mammalia; Eutheria;
                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nature 420:563-573 (2002).
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Rodentia;
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Ludwig W., Gade D., Beck A., Borzym K.,
Schlesner H., Amann R., Reinhardt R.,
"Complete genome sequence of the marine
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01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Sensory transduction histidine kinase.
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                             InterPro;
InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Proc. Natl Acad. Sci. U.S.A. 100:8298-8303 (2003).
-!- SIMILARITY: Contains 1 histidine kinase domain
EMBL; BX294140; CAD73831.1; -.
HSSP: 09ZWQ6; 1JNU.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    strain 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=22735913; PubMed=12835416;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Planctomycetaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacteria; Planctomycetes; Planctomycetacia; Planctomycetales;
                                                                                InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro;
                                                                                                                                                                                                     GO:0011020; C:membrane; IEA.
GO:0001524; F:ATP binding; IEA.
GO:0003577; F:DNA binding; IEA.
GO:00031677; F:kinase activity; IEA.
GO:0016310; F:kinase activity; IEA.
GO:0016740; F:transferase activity; IEA.
GO:0010156; F:two-component response regulator activity; IEA.
GO:0000155; F:two-component sensor molecule activity; IEA.
GO:0000155; F:two-component IEA.
GO:00006315; P:chemotaxis; IEA.
GO:00006160; P:two-component signal transduction system (p. .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; MCI:1930089; MCm3ap.
; MCI:1930089; MCm3ap.
GO:0005737; C:cytoplasm; IDA.
GO:0005634; C:nucleus; IDA.
erPro: IPR005062; SAC3_GANP.
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5; Conserv
p; IPR003594; ATPbind ATPase.
p; IPR000673; CheB methylest.
p; IPR000780; CheR mifrase.
p; IPR011006; CheY like.
p; IPR005467; His Kinase.
p; IPR003661; His kinase.
p; IPR003661; His kinase.
p; IPR003652; N6_Mtase.
p; IPR002052; N6_Mtase.
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1224 AA;
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83.3%;
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Pred. No. 2.1e+03;
1; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOI=10.1073/pnas.1431443100;
Teeling H., Lombardot T.,
m K., Heitmann K., Rabus R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1637
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DR InterPro; IPR000014; PAS.

DR InterPro; IPR000700; PAS-assoc_C.

DR InterPro; IPR001789; Response_reg.

Pfam; PF01339; CheR; 1.

DR Pfam; PF01739; CheR; 1.

DR Pfam; PF02518; HATPase_c; 1.

DR Pfam; PF02518; HATPase_c; 1.

DR Pfam; PF00712; HiskA; 1.

DR Pfam; PF00715; PAC; 3.

DR Pfam; PF00715; PAC; 3.

DR Pfam; PF00072; Response_reg; 1.

PR ProDom; PD000139; Response_reg; 1.

R SMART; SM00387; HATPase_c; 1.

R SMART; SM00388; HiskA; 1.

R SMART; SM00388; HiskA; 1.
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Q6NWV4
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                                                         REPLIENCE FROM N.A.

RESTRAIN=CS7BL/61; TISSUE=Kidney;

REDILINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

REDILINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

REDILINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

REDILINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

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REDILINE=2238257; PubMed=12477932; DOI=10.1073/pnas.242603899;

REDILINE=2238257; PubMed=12477932; DOI=10.1073/pnas.242603899;

REDILINE=2238257; PubMed=12477932; DOI=10.1073/pnas.24260389;

REDILINE=23826032; DOI=10.1073/pnas.24260389;

REDILINE=23826032; DOI=10.1073/pnas.24260389;

REDILINE=23826032; DOI=10.1073/pnas.24260389;

REDILINE=238260389;

REDILINE=2382
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Best Local (
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Fahey J., here
Whiting M., Madar
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcm3ap protein (Fragment).
Name=Mcm3ap;
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05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
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T; SM00138; McTrc; 1.
T; SM00086; PAC; 3.
T; SM00091; PAS; 4.
T; SM000448; REC; 1.
FAM8; TIGR00229; sensory_box; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    472
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PS50123; CHER; 1.
PS50123; CHER; 1.
PS00092; N6 WTASE; U1
PS00092; N6 PAG; 2.
PS50113; PAG; 3.
PS50112; PAG; 3.
PS50110; RESPONSE_REC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VGDIGD 477
                                 Helton E., Ketteman M., Madan A., ., Madan A., Young A.C., Shevchenko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1637
                             Madan A.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                           Chordata;
Rodentia;
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Last sequence update)
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Pred.
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Sciurognathi;
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                                 Shevchenko Y.,
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No.
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has 0; Indels
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thi; Muridae; Murinae; Mus.
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                                 Rodrigues S., Sa
o Y., Bouffard G.
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                                                     Sanchez A.
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RESULT
MC3A_MC
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Best Local
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MGD; MGI:1930089; Mcm3ap.
MGD; MGI:1930089; Mcm3ap.
GO; GO:0005737; C:cytoplasm; ID.
GO; GO:0005634; C:nucleus; IDA.
InterPro; IPR005662; SAC3 GANP.
Dfam: PF03399; SAC3_GANP; 1.
                                                                                                                                                                                                                              for DNA replication.";
Blood 95:2321-2328(2000).
-i- FUNCTION: May be involved in
                                                                                the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed. entitles requires a license agreement (See or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                          Kouno Y., Tom
Sakaguchi N.;
                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N. STRAIN=BALB/C X
                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                      30-MAY-2000 (Rel. 39, Last sequence u5-JUI-2004 (Rel. 44, Last annotation Rel. 44, Last annotation Rokal MCM3-associated protein (GANP Name=Mcm3ap; Synonyms=GANP, Map80;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q9WUU9;
30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jones S.J., Marra M.A.; "Generation and initial analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rodriguez A.C., Grimwood J.,
Krzywinski M.I., Skalska U.,
                                                                                                                                                   This
                                                                                                                                                                                               ++
                                                                                                                                                                                                                                                                  "A novel nuclear phosphoprotein, of the germinal center and associ
                                                                                                                                                                                                                                                                                                     MEDLINE=20197882; PubMed=10733502;
Kuwahara K., Yoshida M., Kondo E., S
Kouno Y., Tomiyasu S., Fujimura S.,
                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MC3A MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=C57BL/6J;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EQUENCE
                                                                                                                                                                        MCM3 (By similarity).
SUBUNIT: Interacts with MCM3.
SUBULIAR LOCATION: Nuclear and cytoplasmic.
PTM: Phosphorylated (Probable).
SIMILARITY: Belongs to the SAC3 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1149
                                                                                                                             SWISS-PROT entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the Engropean Bioinformatics Institute. There are no rest
                                                           AJ006590; CAB44241.1; -.
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    Last sequence update)
    Last annotation updat

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WA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

K Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan R., Farmer A.A., Rubin G.M., Hong L.,

A Hopkins R.F., Jordan R.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

A Hopkins R.F., Jordan R.B., Fonaldo M.F., Casavant T.L., Scheetz T.E.,

A Hopkins R.F., Jordan R.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

A Hopkins R.F., Jordan R., Peters G.J., Abramson R.D., Mullahy S.J.,

A Hopkins R.F., Jordan R., Feters G.J., Abramson R.D., Mullahy S.J.,

A Hopkins R.F., Jordan R., Korrian K.J., Malek J.A., Gunaratne P.H.,

A Hopkins R.F., Jordan R., Jordan R., Jordan R., Garcia A.M., Gay L.J., Hulyk S.W.,

A Hopkins R.F., Jordan R., Young A.C., Shevchenko Y., Gibbs R.A.,

A Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

A Hopkins R.F., Jordan R., Jordan J.W., Green E.D., Dickson M.C.,

Bakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Bakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Bakesley R.W., Touchman J.W., Green E.D., Schein J.S.,

A Karywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

Bakesley R.W., Schein J., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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STRAIN=C57BL/6; TISSUE=Brain;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro;
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                                                                                                                                                                                                                                                                                                                                                                                           mitted (MAY-2003) to the EMBL/GenBank/DDBJ databases.
L; BC052452; AAH52452.1; -.
GO:0005737; C:cyftoplasm; IDA.
GO:0005634; C:nucleus; IDA.
erPro; IPR005062; SAC3 GANP.
m; PF03399; SAC3 GANP; 1.
m; PF03399; SAC3 GANP; 1.
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c. Natl. Acad. Sci. U.S.
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(TrEMBLrel. 26, Last annotation update)
me maintenance protein 3 associated protein.
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Pred. No. 3.5e
1; Mismatches
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Pred. No. 3.5e+03;
1; Mismatches 0;
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Sciurognathi; Muridae;
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RESULT
Q7RNK7
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                                                                                                                               SEQUENCE FROM .....

STRAIN=17XNI;

STRAIN=17XNI;

PubMed=12368865; DOI=10.1038/nature01099;

Carlton J.M., Angiuoli S.V., Suh B.B., Kooij T.W., Pertea M.,

Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L.,

Peterson J.D., Pop M., Kosack D.S., Shumway M.F., Bidwell S.L.,

Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V.,

Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M.,

Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M.,

Florens L., Yates F.R. III, Raine J.D., Sinden R.E., Harris M.A.,

Florens L., Yates F.R., Bergman L.W., Vaidya A.B.,

Gardner M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q80TX5;
01-JUN-2003
01-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Okazaki N., Kikuno R., Ohara R., İnamoto S., Aizawa H., Yuasa S., Makajima D., Nagase T., Ohara O., Koga H.;
"Prediction of the coding sequences of mouse homologues of KIAA gene:
II. The complete nucleotide sequences of 400 mouse KIAA-homologous cDNAs identified by screening of terminal sequences of cDNA clones randomly sampled from size-fractionated libraries.";
DNA Res. 10:35-48(2003).
EMBL, AK122313, BAC65595.1; -.
                                                                                                                 Florens L., Yates F.R. 111, kaine v..., Cunningham D.A., Preiser P.R., Bergman L.W., van Lin L.H., Janse C.J., Waters A.P., Smith van Zalzberg S.L., Venter J.C., Fraser C.M., Hoff
                                                                                                                                                                                                                                                                                                                                                                                                          01-MAR-2004
01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                       Q7RNK7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q7RNK7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MGD; MGI:1930089; Mcm3ap.
GO; GO:0005737; C:cytoplasm; IDA.
GO; GO:0005634; C:nucleus; IDA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Name=Mcm3ap; Synonyms=mKIAA0572;
Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUN-2003 (TrEMBLrel. 24, Creat
01-JUN-2003 (TrEMBLrel. 24, Last
01-MAR-2004 (TrEMBLrel. 26, Last
                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=73239;
                                                                                                                                                                                                                                                                                                                                  Plasmodium yoelii yoelii.
Eukaryota; Alveolata; Api
                                                                                                                                                                                                                                                                                                                                                                   Name=PY01809
                                                                                                                                                                                                                                                                                                                                                                              Hypothetical protein.
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                                                                                "Genome sequence and comparative analysis parasite Plasmodium yoelii yoelii.";
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Pfam; PF03399; SAC3_GANP; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=10090;
             EMBL;
                                                                     Nature 419:512-519(2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=22579291; PubMed=12693553;
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                                       CAUTION: The sequence sh
EMBL/GenBank/DDBJ whole
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            preliminary data.
; AABL01000489; EAA21180.1;
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                                                                                                                                                                                                                                                                                                                                     Apicomplexa; Haemosporida; Plasmodium.
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                                                        shown here
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Pred. No. 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C980215AE6B2D440 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence update)
                                         shotgun
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                                                        is derived from
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DR InterPro; IPR006209; EGF_like.

DR PROSITE; PS00022; EGF_1; UNKNOWN 1.

DR PROSITE; PS00022; EGF_1; UNKNOWN_I.

KW Hypothetical protein.

SQ SEQUENCE 2251 AA; 267817 MW; F3B6FCB24BBD5492 CRC64;

Query Match
Best Local Similarity 83.3%; Pred. No. 4e+03;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Matches 5; Conservative 1; Mismatches 0; Todels 0; Saps 0;

Search completed: July 27, 2005, 01:20:29

Job time: 178 secs
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